

# STIC Search Report Biotech-Chem Library

#### STIC Database Tracking Number

TO: Andrew D Kosar Location: 3c04 / 3c18

Art Unit: 1654

Wednesday, May 10, 2006

Case Serial Number: 10/800179

From: Noble Jarrell

**Location: Biotech-Chem Library** 

**Rem 1B71** 

Phone: 272-2556

Noble.jarrell@uspto.gov

Search Notes		·	
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#### STIC-Biotech/ChemLib

From: Sent: To: Subject:	Tuesday, May 09, 20 STIC-Biotech/Chemi		
Requester:  ANDREW KOSAF Art Unit:  GROUP ART UN Employee Number:  80341 Office Location:  REM 03C04 Phone Number:  (571)272-091 Mailbox Number:  REM 3c18  Case serial number  10/800,179	NIT 1654 13		
Class / Subclass ( Earliest Priority			
Format preferred Paper Search Topic Info	ormation:	.9 in 10/800,179.	
Special Instructi			
*****************  Searcher:	6	*************  Type of Search  NA# AA#:  S/L: Oligomer:  Encode/Transl:  Structure #: Text:  Inventor: Litigation:	********* Vendors and cost where applicable STN: DIALOG: QUESTEL/ORBIT: LEXIS/NEXIS:_ SEQUENCE SYSTEM: Compug WWW/Internet: Other (Specify):

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# PALM INTRANET

Day: Wednesday Date: 5/10/2006 Time: 16:58:59

# **Inventor Information for 10/800179**

Inventor Name	City	State/Country
KUMAR, MANOJ	FREMONT	CALIFORNIA
CUEVAS, WILLIAM A.	SAN FRANCISCO	CALIFORNIA
Appln Info Contents Petition II	nfo Atty/Agent Info Contin	nuity Data Foreign Data Inve
Search Another: Application#	Search or Patent#	¥ Search
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To go back use Back button on your browser toolbar.

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### **Protein Sequence Searches - February 2005**

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

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#### November 2005

Published\_Applications Nucleic Acid and Published\_Applications Amino Acid database searches now generate two sets of results each. The Published\_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Newly published applications will appear in the Published\_Applications\_New databases; older published applications make up the Published\_Applications\_Main databases.

Searches run against Nucleic Acid Published\_Applications produce two sets of results, with the extensions .rnpbm (Published\_Applications\_NA\_Main) and .rnpbn (Published\_Applications\_NA\_New).

Searches run against Amino Acid Published\_Applications produce two sets of results, with the extensions .rapbm (Published\_Applications\_AA\_Main) and .rapbn (Published\_Applications\_AA\_New).

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elastin precursor - mouse

N;Alternate names: tropoclastin
C;Species: Mus musculus (house mouse)
C;Date: 18-Aug-1995 #sequence_revision 16-Aug-1996 #text_chan
C;Accession: A55721
R;Wydner; K.S.; Sechler, J.L.; Boyd, C.D.; Passmore, H.C.
Genomics 23, 125-131, 1994
A;Title: Use of an intron length polymorphism to localize the
A;Reference number: A55721; MUID:95130069; PMID:7829060
A;Accession: A55721
A;Molecule type: mRNA
A;Residues: 1-860 <WYD>
A;Cross-references: UNIPROT:P54320; UNIPARC:UPI0000029BE6; GB
C;Genetics:
A;Map position: 5
C;Superfamily: elastin
C;Keywords: alternative splicing; extracellular matrix; glyco
F;1-27/Domain: signal sequence #status predicted <MATI>
F;850-855/Disulfide bonds: #status predicted
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                                                                                                                                                    G-PGVGPGVG-PGAGAGS---GAGAGSGA---GAGSGAGAGSGVGVP-GVGVPGVGVPGKG
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                                                                                                                                                                                                                                                                                                                                                                 AAAAKAAKYGAGGAGV-LPGVG--GGGIPG----GAGAIPGIGGIAGAGTPAAAAAAAKA
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                                                                                     ---PAAAAAAAKAAAKAAQYGLG-GAG-
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Pred. No. 1.2e-66;
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                                           -GLGAG-GLGAGGGVSPAAAAKAAKYGAAGLG
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RESULT 3
EART
elastin precursor - rat
N;Alternate names: tropoelastin
C;Species: Rattus norvegicus (Norway rat)
C;Date: 11-Jan-1991 #sequence revision 16-Aug-1996 #text\_change 09-Jul-2004
C;Accession: A36106; A30878; \overline{A}36523; S03173; I54172; I68505
R;Pierce, R.A.; Deak, S.B.; Stolle, C.A.; Boyd, C.D.

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Biochemistry 29, 9677-9632, 1990
A.Field before nambgrrily of rat tropopolastin mRNA revealed by cDNA cloning.
A.Field before nambgrrily of rat tropopolastin mRNA revealed by cDNA cloning.
A.Field before nambgrrily of rat tropopolastin mRNA revealed by cDNA cloning.
A.Field before nambgrrily of rat tropopolastin is synthesized from a 3.5-kilobase mRNA.
A.Field before nambgrrily 1800-1807, 1889
A.Fille Rat tropopolastin is synthesized from a 3.5-kilobase mRNA.
A.Field before nambgrrily is synthesized from a 3.5-kilobase mRNA.
A.Field before nambgrrily is synthesized from a 3.5-kilobase mRNA.
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A.Field before nambgrrily is synthesized from a 3.5-kilobase mRNA.
A.Field before nambgrrily is synthesized mRNA.
A.Field before nambgrrily is synthesized from a 3.5-kilobase mRNA.
A.Field before nambgrrily is synthesized mRNA.
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140
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                                                                                                                                                                                                                                                                                                                                                             n 28.4%; Score 1185; DB 1;
Similarity 43.8%; Pred. No. 1.1e-64;
78; Conservative 47; Mismatches 299;
                                                         GPGA-GAGSGAGAGSGAGAGSGAGAGSGVGVP--GVGVPGVGVPGKGVPGVG-PGVGP-G 161
                                                                                                                                                                                                                                          GVPGAVPGGVPG-GLPG-GVPGGVYYPGAGIGGGLGGGALGPGGKPPKPGAGLLGAFGAG
                                                                                                                                                                                                                                                                                             GVPGV---GVPGVGVPGKGVPG----VGPGVGPGVGPG-VGPGA-----GAG----SGAG
VPGAVGVGGVPGAVGGIGGIGGLGVSTGAVVPQLGAGVGAGGKPGK-VPGVGLPGVYPGG
                                                                                                                  PGGLGGAGPGAGLSYASRPGGVLVPGGGAGAAAAYKAAAKAGAGLGGIG-GVPGGVGVGG 139
                                                                                                                                                                           AGSGAGAGSGAGAGSGVGVPGVGVPGVGV-----PGKGVPGVGPGVGPGVGPGV 106
                                                                                                                                                                                                                                                                                                                                                                                                                         Length 864;
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198
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BABO

elastin precursor, splice form a - bovine

N.Alternate names: tropoelastin

N.Contains: elastin precursor, splice form b; elastin precursor,

C.Species: Bos primigenius taurus (cattle)

C.Date: 08-Jun-1999 #sequence revision 26-Jul-1996 #text change C

C.Accession: A31865; A26728; B26728; C26728; A22343; I45886

C.Accession: A31865, A26728; B26728; C26728; A22343; I45886

R.Yeh, H.; Anderson, N.; Ornstein-Goldstein, N.; Bashir, M.M.; RC
                                                           A;Cross-references: UNIPROT:P04985; UNIPROT:Q28101; UNIPARC:UPI000016C304; GB:J02855; R;Raju, K.; Anwar, R.A.
J. Biol. Chem. 262, 5755-5762, 1987
                                                                                                                                                                                    Biochemistry 28, 2365-2370, 1989
A;Title: Structure of the bovine elastin gene and S1 nuclease A;Reference number: A31865; MUID:89274159; PMID:2543440
J. Biol. Chem. 262, 5755-5762, 1987
A;Title: Primary structures of bovine elastin a, b, and A;Reference number: A92640; MUID:87194772; PMID:3032943
A;Accession: A26728
                                                                                                                        A; Molecule type: DNA
A; Residues: 1-27 < YEH>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGGAG-GLGVGGKPPKPYGGALGA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGSGAGAGSGAGAGS----GAGAGSGVGVPGVGVPGVGVPGVGPGVGPGVGPGVG 587
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                                           ведчелсев
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A;Molecule type: mRNA A;Residues: 1,'RS',4-11,'E',13-636,'V',638-747 <RAJ> A;Cross-references: UNIPARC:UPI0000129E6C; GB:J02717;

NID:g163019; PIDN:AAA30503.1;

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A;Introns: 634/3; 653/3; 676/3; 689/3; 707/3; 716/3; 733/3
A,Note: the list of introns is incomplete
C;Superfamily: elastin
C;Keywords: alternative splicing; extracellular matrix; glycoprotein; hydroxylysine
F;1-747/Product: elastin precursor, splice form a #status predicted <EPA>
F;1-225,260-747/Product: elastin precursor, splice form b #status predicted <EPB>
F;1-225,240-747/Product: elastin precursor, splice form b #status predicted <EPB>
F;1-26/Domain: signal sequence #status predicted <EPB>
F;27-747/Product: elastin #status predicted <MAT>
F;27-747/Product: elastin #status predicted <MAT>
F;105,109,252,271,275,324,327,400,404,407,445,448,489,493,544,548,552,606,609,645,64
F;737-742/Disulfide bonds: #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule type: DNA
A;Residues: 678-683,685-747 <ROS>
A;Residues: 678-683,685-747 <ROS>
A;Residues: 678-683,685-747 <ROS>
A;Residues: 678-683,685-747 <ROS>
A;Residues: 678-683,685-747 <ROS
A;Conserve Feferences: UNIPARC:UPI000008856A; GB:M31898; NID:g163015; PIDN:AAA96417.1;
R,Brown, P.L.; Mecham, L.; Tisdale, C.; Mecham, R.P.
Blochem. Biophys. Res. Commun. 186, 549-555, 1992
Blochem. Biophys. Res. Commun. 186, 549-555, 1992
A;Title: The cysteine residues in the carboxy terminal domain of tropoelastin form a A;Reference number: A58621, MUID:92337651; PMID:1632791
A;Contents: annotation, disulfide bonds
C;Comment: The term tropoelastin refers to a soluble precursor form of the extracell
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Lab. Invest. 51, 605-623, 1984
A;Title: Biology of disease: Elastin: Relation of protein
A;Reference number: I45885; MUID:85059254; PMID:6150137
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A;Cross-references: UNIPARC:UPI000002A8D4; GB:K03506; NID:91630;
R;Cicila, G.; May, M.; Ornstein-Coldstein, N.; Indik, Z.; Morrov
Biochemistry 24, 3075-3080, 1985
B;Title: Structure of the 3' portion of the bovine elastin gene
A;Reference number: A22343; MUID:85280426; PMID:2992576
A;Accession: A22343
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A;Molecule type: mRNA
A;Residues: 1,'RS',4-11,'E',13-225,240-636,'V',638-747 <RA2>
A;Cross-references: UNIPARC:UPI000002A8D3; GB:K03505; NID:g163025; PIDN:AAA30505.1;
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A; Residues: 613-747
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Best Local Similarity
Matches 350; Conserv
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                               GPGIPG--
                                                                               PTGAGVKPKAQVGAGAFA-----GIPGVGPFGGQQPGLPLGYPIKAPKLPAGYGL
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                            -APGAIPGIGGIAGVGAPDAAAAAAAAAAKAAKFGAAGGLPGVGVPGVGVP
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41.4%; Pred. No. 1.1e-62;
tive 46; Mismatches 249)
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RESULT 5

S59623

S59623

troppelastin - sheep

C;Species: Ovis orientalis aries, Ovis ammon aries (don C;Species: Ovis orientalis aries)
C;Date: 23-Aug-1996 #sequence_revision 13-Mar-1997 #tep
C;Accession: S59623, A24758
R;Mauch, J.C.; Sandberg, L.B.; Roos, P.J.; Jimenez, F.;
RMartix Biol. 14, 635-641, 1994
A;Title: Extensive alternate exon usage at the 5' end of A;Reference number: S59623
A;Accession: S59623
                                                                                                                                       A;Cross-references: UNIPROT:P11547; UNIPARC:UPI0000177409
R;Yoon, K.; Davidson, J.M.; Boyd, C.; May, M.; LuValle, P.; Ornstein-Goldstein, N.;
Arch. Blochem. Biophys. 241, 684-691, 1985
A;Title: Analysis of the 3' region of the sheep elastin gene.
A;Reference number: A24758; MUID:85305763; PMID:3839997
A;Accession: A24758; MUID:85305763; PMID:3839997
A;Accession: A24758
A;Hosidues: 655-669,671-716,732-770 <YOO>
A;Gross-references: UNIPARC:UPI0000129E70
C;Reywords: alternative splicing; extracellular matrix; glycoprotein; hydroxylysine F;760-765/Disulfide bonds: #status predicted
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A; Residues: 1-770 < MAU>
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                                                                       Query Match
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Matches 368
                                                                                            Local Similarity
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 GGVLGAVPG-GVPGGVFFPGAGLGGLGVGGLGPGVKPAKPGVGGLVGPGLGAEGSGLLPA
                                     GKSCG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         --AAAAAAKAAAKAAQFGLG-GVGGLGVGGLGAVPG----AVGLG----GVSPAAAAKAAK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -----AAKAAKIGAGGVGALGGVVP--GAPGAIPGL-PGVGGVPGVGIPAAAAAKAAA
                                                                         Conservative
                                                                                                     27.5%;
                                                                                          42.5%;
                                                                           41; Mismatches
                                                                                          Score 1146.5; DB Pred. No. 2.1e-62;
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-GAGSGAG---

AGSGAGAGSGVGVPGVGVPGVGVPGVGPGVGPG

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Smit)

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A;Molecule type: DNA
A;Residues: 1-713 <SH>
A;Residues: 1-713 <SH>
A;Residues: 1-713 <SH>
A;Cross-references: UNIPROT:P08399; UNIPARC:UPI000016D179; GB:X02966; GB:M12039; NID:955
C;Comment: Mutations within the per locus of the fruit fly affect a variety of natural h
logous locus with multiple tandem repeats of nucleic acid hexamers (ACNGGN, TCAGGC) that
C;Comment: The serine residues of the S-G repeats found in certain proteoglycans are atb
                                                                                                   period clock protein - mouse (fragment)
(;Species: Mus musculus (house mouse)
(;Species: Mus musculus (house mouse)
(;Date: 31-Mar-1988 #sequence_revision 31-Mar-1988 #text_change 09-Jul-2004
(;Accession: A24403
R;Shin, H.S.; Bargiello, T.A.; Clark, B.T.; Jackson, F.R.; Young, M.W.
R;Shin, H.S.; Bargiello, T.A.; Clark, B.T.; Jackson, F.R.; Young, M.W.
Nature 317, 445-448, 1985
A;Title: An unusual coding sequence from a Drosophila clock gene is conserve
A;Reference number: A24403; MUID:86014384; PMID:2413365
A;Accession: A24403
                                                                                                                                                                                                                                                         RESULT
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                                                                                                                                                                                                                                                                                                                                                                                         VSPAAAAKAAKFGAAGLGGVLGAGRPFPIGGVAARPGFGLSPIFPG-GAGGLGVGGKPPK
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C;Keywords: circadian rhythm; tandem
F;41-77/Domain: EGF homology <EGF>
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                                                 GTGTGTGTGTGTSTVTVRGTGT---
                                                                         GSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGSGAGA
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nilarity 35.3%;
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Pred. No. 7.2e-62;
4; Mismatches 304
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RESULT 7

F70806

hypothetical glycine-rich protein Rv3508 - Mycobacterium tuberculosis (strain H Protein Ry508)

C;Species: Mycobacterium tuberculosis

C;Species: Mycobacterium tuberculosis

C;Date: 17-Jul-1998 #sequence\_revision 17-Jul-1998 #text\_change 09-Jul-2004

C;Accession: F70806

R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.;

C;Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Ho

Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S.

Nature 393, 537-544, 1998

A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G.

A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complet complete Holroyd, H37RV) Gordon, genome s s

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A;Reference number: A70500; MUID:98295987; PMID:9634230
A;Accession: F70806
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-1901 <COL>
A;CCOSE-references: UNLIPROT:053553; UNIPARC:UPI000013C2A6; GB:AL022022; GB:AL123456;
A;Experimental source: strain H37Rv
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C;Superfamily:
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                                        GLS-GFDGGQGGQGGAGGSAGAGGINGAGGAGGTGGAGGDGAPATLIGGPDGGDGGQGGI
                                                                GVGPGVGPGVGPGAGAGSGAGAG--SGAGAGSGAGAGSGVGVPG---------
                                                                                                                       AGGAGGAGDNNFNGGOGGAGGOGGLGGASTTS INANGGAGGNGGTGGKGGAGGAGTL
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Pred. No. 4.3e-59;
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RESULT 8
B70807
C;Species: Mycobacterium tuberculosis
C;Species: Mycobacterium tuberculosis
C;Species: Mycobacterium tuberculosis
C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 20-Jun-2000
C;Date: 17-Jul-1998 #sequence, T.; Garnier, T.; Churcher, C.; Harris, D.; Gorder, Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroy, Connor, R.; Davies, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G.
A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G.
A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete get A;Reference number: A70500; MUID:98295987; PMID:9634230
A;Reference number: A70500; MUID:98295987; PMID:9634230
A;Residues: 1-1079 < COL>
A;Coss-references: UNIPARC:UPI00000D3AEA; GB:AL123456; NID:g3261554; A;Gene: Rv3512
C;Genetics:
A;Gene: Rv3512
C;Superfamily: collagen alpha 1(I) chain; fibrillar collagen carboxyl-terminal homo:
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GTGGQGGAGGDGGNGAIGGHGPLTDDPGGNGGTGGNGGTGGTGGAGIGSLGGGTGGDGGN
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                                                  -----GSGAGAGSGA---
                                                                                                     GTGGTGGSGGTGGDGGNGADNTANMTAQAGGDGGNGGDGGFGGGAGAGGGGLTAGAN
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                                                  GAGSGVGVPGVGVPGVGVPGVGPG
                                                                                                                                                     -GPGVGPGVGPGAGAGSG---AGA-
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R; Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gor, Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holro Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998

A; Authors: Squares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A; Title: Deciphering the biology of Mycobacterium tuberculosis from the complete g. A; Reference number: A70500; MUID:98295987; PMID:9634230

A; Resession: D70807

A; Status: preliminary; nucleic acid sequence not shown; translation not shown A; Molecule type: DNA

A; Cross-references: UNIPARC:UPI00000D3AEB; GB:AL022022; GB:AL123456; NID:93261554; A; Cross-references: Strain H37Rv
C; Gene: Rv3514
C; Superfamily: collagen alpha 1(IV) chain
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C;Species: Mycobacterium tuberculosis
C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 20-Jun-2000
C;Accession: D70807
C;Accession: D70807
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.;
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.;
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                                                                                                                                                                                                                                                                                                                      VGVPGVGVPG--VGVPGK-GVPGVGPGVGPGVGPGVG-----PGAGAGSGAGA----
                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGGAAGTGGTGGMIGTTGNAGVGGAG---GQGGDGGAGGAGADADQPGATGGTGFAGGAGG
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  GGTGGVGGTGGDGGNAGTGAGDPGKG-GTGGTGGTGGSGGAGGSGGANFNGGTGGTGGTG
                                                                                        DQPGATGGTGFAGGAGGAGGSGGSSCAGGTNGSGGAGGTCGQVVAGGAGISFSNGSNGGT
                                                                                                                                                                                GIGGTGGDGGTGGAAGAGGAAGTGGTGGMIGTTGNAGVGGAGGQGGDGGAGAGADA
                                                                                                                                                                                                                           ----GSGAGAGSGAGAGSGVGVPGVFGVPGV--VGVPGK-GVPGVGPGVGPGVGPGVGPGVG
                                                                                                                                                                                                                                                                        GGAAGTGGTGGMIGTTGNAGVGGAGGSSGAGGTNGSGGAGGTDGQGGAGGAGADNPT
                                                                                                                                                                                                                                                                                                                                                                                                                ------GSGAGAGSG
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                                                                                                                                     PGAGAGSGAGAGSGAGAGSG----
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                                          GVPGKGVPGVGPGVG---PGVGPGVGPGAGAGSGAGAGSGAGA----GSGAGAGSG 313
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36.0%;
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                                                                                                                                     ---AGAGSGVGVPGVGVPGV-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VGPGVG-PGVGPGVGPGAGAGSGAGAGSG------AGAGSGAGAGAGSGVGVPGVGV 681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPG--VGVPGKGVPGVGPGVGPGV--GP
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    AATGVGDGGDGGNG-GNGGNGGTGVGSPGGLGGAGGTGGLGGAGAGGGA
                                                                                        AGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGAGAGSGAGA
                                                                                                                                                                                                                                                                                                                                                                                 AGTGAGDPGKGGTGGTGGSGGAGGSGGANFNGGTGGTGGTGGKGGMGGIAGDGG 1366
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGGQCGAGGAG I S F S NG S NG G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G T G G 
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                                                                                                                                                                                                                                                                                   PG----VGVPGK-----GVPGVGPGVGPGVGPGVGPGAGAGSGAGAGSGAGAGSGAG
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A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-453,483-617,651-792 <FAZ>
A;Residues: 1-453,483-617,651-792 <FAZ>
A;Residues: 1-453,483-617,651-792 <FAZ>
A;Residues: 1-453,483-617,651-792 <FAZ>
A;Residues: UNIPARC:UPIO000129E6E; EMBL:M36860; NID:g182061;
A;Note: this sequence represents a composite of several splice forms
R;Fazio, M.J.; Olsen, D.R.; Kuivaniemi, H.; Chu, M.L.; Davidson, J.M.
Lab. Invest. 58, 270-277, 1988
A;Title: Isolation and characterization of human elastin cDNAs, and a
A;Reference number: A53891; MUID:88156138; PMID:2831431 A;Cross references: UNIDARC:UPI00006E12E; GB:J04821; NID:g182052; R;Fazio, M.J.; Olsen, D.R.; Kauh, E.A.; Baldwin, C.T.; Indik, Z.; O J. Invest. Dermatol. 91, 458-464, 1988
A;Title: Cloning of full-length elastin cDNAs from a human skin fib A;Reference number: A30524; MUID:89009960; PMID:3171221
A;Accession: A30524 elastin precursor, long splice form N;Alternate names: tropoelastin C;Species: Homo sapiens (man) C;Date: 22-Uun-1990 #sequence revisi C;Accession: A32707; A33705; A30524; R;Bashir, M.M.; Indik, Z.; Yeh, H.; Ornstein-Goldstein, J. Biol. Chem. 264, 8887-8891, 1989
A;Title: Characterization of the complete human elastin A;Reference number: A33705; MUID:89255358; PMID:2722804 A;Molecule type: mRNA
A;Residues: 1-500,507-792 <IND>
A;Cross-references: UNIPROT:P15502; R;Indik, Z.; Yeh, H.; Ornstein-Goldstein, N.; Sheppard, P.; Ar Proc. Natl. Acad. Sci. U.S.A. 84, 580-5864, 1987 A;Title: Alternative splicing of human elastin mRNA indicated A;Reference number: A32707; MUID:87289668; PMID:3039501 A; Molecule type: DNA A; Residues: 1-27 <BAS> A; Accession: A33705 A; Accession: A32707 ;Species: Homo sapiens (man)
;Date: 22-Jun-1990 #sequence revision 26-Jul-1996 #text\_change 09-Jul-2004
;Accession: A32707; A33705; A30524; A53891 10 UNIPROT:Q9UMK5; UNIPARC:UPI0000173C4E; GB:M16983; Ornstein-Goldstein, N.; Rosenbloom, J.C.; Abrams, P.; Anderson, à Delineation ۲. ۲. sequence Ornstein-Goldstein, N. age-associated PIDN: AAA52382.1; N.; Rosenbloom, analysis of cl of unusual

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A;Cross-references: GDB:119107; OMIM:130160
A;Map position: 7q11.23-7q11.23
C;Superfamily: elastin elastin; extracellular matrix; C;Keywords: alternative splicing; extracellular matrix; F;1-26/Domain: signal sequence #status predicted <AMI>F;27-792/Product: elastin #status predicte
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A;Accession: A53891
A;Molecule type: mENA
A;Molecule type: mENA
A;Residues: 164-453,483-500,507-617,651-792 <FA2>
A;Cross-references: UNIPARC:UPI000016A887; GB:M24782; NID:g182063; PIDN:AAA53190.1; PID:
A;Cross-references: UNIPARC:UPI000016A887; GB:M24782; NID:g182063; PIDN:AAA53190.1; PID:
C;Comment: The term tropoelastin refers to a soluble precursor form of the extracellular
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Matches 335
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                                          VPGALAAAKAAKYGAAVPGVLGGLGALGGVGI-PGGVVGAGPAAAAAAAKAAAKAAQFGL
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40.6%; Pred. No. 6e-5
tive 47; Mismatches
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Db 710 VGAAGLG--GLGVGGLGVPGVG-GLG-GIPPAAAAKAAKYGAAGLG 751
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major ampullate fibroin protein - orb spider (Nephila clavipes) (fragment) (C;Species: Nephila clavipes C;Date: 08-Mar-1991 #sequence\_revision 13-Jan-1993 #text\_change 09-Jul-2004 C;Accession: A36068 R;Xu, M.; Lewis, R.V. Proc. Natl. Acad. Sci. U.S.A. 87, 7120-7124, 1990 Proc. Natl. Acad. Sci. U.S.A. 87, 7120-7124, 1990 A;Title: Structure of a protein superfiber: spider dragline silk. A;Reference number: A36068; MUID:90384959; PMID:2402494 A;Accession: A36068 A;Status: preliminary A;Accession: A36068 A;Status: preliminary A;Molecule type: DNA A;Mole A; Residues:  $\bar{1}$ -718 < XUA> A; Cross-references: UNIPROT:P19837; UNIPARC:UPI000016BF56; GB:M37137; A; Note: the authors translated the codon GGT for residue 292 as Gln, G 121 GAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS 48 AAAAG---GAGQG---61 GAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS 1 GAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS GAGAGSGAGAGSGVGVPGVGVPGVGVPGVGVPGVGPGVGPGVGPGVG-PGAGAGSGAGAG YGGLG-GQGAGQG-GYGGLGSQGAGRGGLGGQGAGAAAAAAAAG-GAGQGGLGGQGAGQGA GAGAGSGAGAGSGVGVPGVGVPGVGVPGVGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS GAGAGSGAGAGSGVGVPGVGVPGVGVPGVGVPGVGPGVGPGVGPGAGAGSGAGAGS AGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSG SGAGAGSGAG----AGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSG GAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS AAAGGAGQGGYGGLGSQGAGRGGLG--GQGAGAVAAAAAGGAGQG-GYGGLGSQGAGRG-GAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS YGGLG-GQGAGQG-GYGGLGSQGAGRGCGLGGQGAGAAAAGGAGQGGLGGQGAGQAAAAA GAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS GASAAAAGGAGQG-----GAGAGSGAGAGSGVGVPGVGVPGVGVPGVGVGVGVGPGVGPGVGPGAGAGSGAGAGS AAAGGAGQGGYGGLGSQGAGRGGLGGQGAGAAAAAAAAGGAGQG-GYGGLGNQGAGRG-GQ GAGAAAAAAAGGAGQG----GYGGLG-----GQGAGQGGYGGLG-GQGAGQGAGAAA AAAAG---GAGQG-GYGGLGNQGAGRGGQGAAAAAAGGAGQGGYGGLGSQGAGRG-GQGAG GQGAGAAAAAAGGAGQRGYG--GLGNQGAGRGGLG--GAAAAAAGGAGQG-----GYGGLGSQGAGRGGLG-----AGQGGLGGQGAGQGAGAAAAAAG-GVRQGGYGGLGSQGAGRG-Conservative 38.4%; 24.9%; 39; -----GYGGLGSQGAGRG Score 1041; DB 2; Pred. No. 4.1e-56; Mismatches -GYGGLGSQGAGRG-----GEGAGAAAAAAGGAGQGG DB 2; 305; Length 718; Indels -GQGAGAAAAAAGGAGQGG 138; *YGAAAAAAAGG* GOGAGAAA GQGAGAAA ; NID:g159711; PID:g1 GTA for residue 374 Gaps ဥ္ပ 589 715 545 655 494 599 439 540 398 480 342 420 284 360 240 300 183 240 136 180 78 120 47 60 32

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RESULT 12
A70869
hypothetical glycine-rich protein Rv2490c - Mycobacterium tuberculosis (strain H37RV)
C.Species: Mycobacterium tuberculosis
C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 09-Jul-2004
C;Accession: A70869
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gordon, S.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroyd, S. Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S.
Nature 393, 537-544, 1998
A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G.
A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete genome A;Reference number: A70500; MUID:98295987; PMID:9634230
A;Recession: A70869
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Cross-references: UNIPROT:053215; UNIPARC:UPI00001652F8; GB:AL021246; GB:AL123456; NIC A;Experimental source: strain H37Rv
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A;Gene: Rv2490c
C;Superfamily:
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                                 GAGGDGGDPGAG----GKGGAGGAGATEGVTGATGATVHSGGNGGKGGNGADATVAGANGG
                                                                  GKGVPGVGPGVGPGVGPGAGA-----
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-SGAGAGSGVGVPGVGVPGVGVPG-----KGVPGVGP-----GVGP
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                                                                  --GSGAGAGSGA-----
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hypothetical glycine-rich protein Rv1450c - Mycobacterium tuberculosis (strain H37RV) C;Species: Mycobacterium tuberculosis C;Species: Mycobacterium tuberculosis C;Date: 17-Jul-1998 #sequence\_revision 17-Jul-1998 #text\_change 20-Jun-2000 C;Accession: E70917

R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gordon, C.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroyd, S.; Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998

A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete genome A;Reference number: A70500; MUID:98295987; PMID:9634230

A;Accession: E70917

A;Status: preliminary; nucleic acid sequence not shown; translation not shown တ တ

GB:AL123456;

NID:g3250713;

PIDN:CA

genome

A;Molecule type: DNA
A;Residues: 1-1329 <COL>
A;Cross-references: UNIPARC:UPI00000C158B; GB:Z95844;
A;Experimental source: strain H37Rv
C;Genetics:
A;Gene: Rv1450c
C;Superfamily: collagen alpha 2(I) chain; fibrillar co collagen carboxyl-terminal homology

Query Match Local 134 291 177 al Similarity 33.8 340; Conservative 88 41 GVGPGVGPGAGAGS-----GAGAGSGA-----GHGAEGVAGAAGEDLSPHGTSGGVGGDAGDGGTGGRGGWLAGAGGAGGAGGVGGTGGAGG GKGVPGVGPGVGPGVGP-----GVGPGAGAG-----SGAGAGSGAGAGSGAGAGSG 133 G---GVGGTGGAGGLGGTLYGAGGHGGAGGPGPIGGVGGHGGVGGAAGLLGVGGHG-GAG ĠĀĠĀĀGĠĀĠGSGĠWLLĠNGĠVĠĢĀĠĢQSLLGGĀTGGĀGGNĀGLFĠVĠĠTĠĢÞĠ-ĠÞĠ-ĠÞ GAGAGSGAGAGSG--VGVPGVGVPG-----VG-----VPG--VGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGA-----GS 180 24.2%; Score 1010.5; DB 2; 33.8%; Pred. No. 4.4e-54; ative 35; Mismatches 382; -----VGVPGKGVPGVGPGVGP -GAGSGAGAGSGVGVPGVGVPGVGVP Indels 249; Length 1329; Gaps 350 290 87 234 40 47

181 GAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS 240

AGFSRALIVAGDNGGDPGAGGAG-GTGGAGSTIGAHGAAGASPTSGGNGGAGGNGAHFSS

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hypothetical glycine-rich protein Rv3507 - Mycobacterium tuberculosis (strain H37Rv C;Species: Mycobacterium tuberculosis C;Species: Mycobacterium tuberculosis C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 20-Jun-2000 C;Accession: E70806 R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gord, Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroy Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998 A;Authors: Sqares, R.; Sulston, J.B.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete ge A;Reference number: A70500; MUID:98295987; PMID:9634230 A;Accession: E70806 A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Residues: 1-1381 cCOL> A;Cross-references: UNIPARC:UPI0000D3AE7; GB:AL022022; GB:AL123456; NID:g3261554; A;Cross-references: UNIPARC:UPI0000D3AE7; GB:AL022022; GB:AL123456; NID:g3261554; A;Genetics: Rv3507 C;Superfamily: collagen alpha 1(IV) chain
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TGGNAGSQGIG-----GDGGNGGDGGKGGDGADA
                                                                                                                                                                                                                                           GGNAGGAGGQGGNAGQGGAGGNAVIPGDGV-----GKAPHGDAGGSGGDGGKGGQG
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                             VPGK-GVPGVGPGVGPGVGPGAGAGSGAGA
                                                                                      PGVGPGVGPGAGAGSGAGAGSGA------GAGSGAGAGSGVGVPG-VGVPGVG
                                                                                                                     GNGGNAGAGGNGGSGDFGGNTTSGASGSGGNGGNAGTAGSG----GAGGTGGTGLSGGNG
                                                                                                                                                 GPGAGAGSGAGAGSG-----AGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVG
                                                                                                                                                                                  GSGGTGGSGAPIGGGAGGTGGSGGHAGKGGAG-
                                                                                                                                                                                                             AAGANGGTGGSGGNGGDGGAGGTGAAG--GNGTPGTGPAGGTGAKGGDGGDGGAG-GA
                                                                                                                                                                                                                                                                                                                                                              NGGTAGSQGTGGVGGDGGAGGNGGGGKAGTGNSGNFGVDGE----AGFSGGAGGNGGVGG
                                                                                                                                                                                                                                                                                                                                                                                        SGAGAGS----GAGAGSGAGAGSGVGVPGVGVPG-VGVPGKGVPGVGPGVGPGVGPGVGP
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                                                         GNGGNG-GNGGDGGNGAHGTVGAQFVPATSLPTPNGGAGGNGGTGSNGGAPGPAGAPGPT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24.1%; Score 1006; DB 2; 35.1%; Pred. No. 8.4e-54; tive 31; Mismatches 384;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -GVGPGVGPG----AGAGSGAGAGSGAGAGSGAGA-----GSGV
                                                                                                                                                                                  -GIGAQGTTITVPGNGGNAGDG-----
                             778
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RESULT 15
A70934
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, I Mycobacterium tuberculosis (strain D.; Gordon, တ

695	GPGVGPG-VGPGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVG	Оу 639
954	GGNGGNGGNGAAGGNGGNGGNGGNGSLRGSGAGGHGGNG-GNASRGMGGDGGTG	Db 896
638	GVEGVEGVEGVEGVEGVEGVEGVEGVEGVEGVEGVEGVEG	Оу 599
895	PGDGGNGANGSVFTNNGIGGNG GNAGCNAGPSGAGGSGGAGSTFGATGSSSSSIHVN	Db 841
598	acscvgvpqvgvpqvgvpqxqvpqqvqpqqpqpqpqqqpqqqqqqqqqq	Qy 550
840		Db 788
549	GVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGSGAGAGSGAG	Оу 500
787	GHPGGTGGVGGTGGKGGDGGDGGAAPNGVAGSQGPGGAGGDGGTGGVGGNGGRGIDGADG	Db 728
499	GVPGKGVPGVGPGVGPGVGPGVGPGAGAGS-GAGAGSGAGAGSGAGAGSGVGVPGV	Оу 445
727	NGSSA	σ,
444	VGPGVGPGVGPGVGPGAGAGSGAGAGSGAGAGSGAGAGSGV-GVPGVGVPGV	Оу 394
667		0
393	GSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPG	Оу 359
607	AGGHGGD	
358	AGAGSGVGVPGVGVPGVPG-KGVPGVGPGVGPGVGPGVGPGAGAGSGAGA	Оу 308
549	GDGGSGGAGSIGFNASAPGAAGSPGGNGGNGGPGGAGGEGGAGGLALAAASGQNGSQG	Db 493
492	Ċ	Dv 252
251	GVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGSGAGAGSGAGAG	
434	GAPGTTS	
199	GVGPGVGPGVGPGAGAGSGAGAGSGVGVPGV	
374	VGGPGGGHGL-LGLGGSHGAGGAGGSGGDGGAPGDGGNGATGTWGHNLGAGGTGGNGGNP	Db 316
156	GAGSGAGAGSGVGVPGVGVPG	Оу 97
315	GAGGAGGVLMGNGGNGGAGGEGGPGGAGGAGGAGGAHATNI.GADGQAGGNGGNGGNGGAGGTGG	Db 256
96	GAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGP	Qy 51
255	GAGA-VGGNGGAGGTAGLFGVGGAG-GAGGNGIAGVTGTSASTPGGSGTAGGAG-GIGGNG	Db 198
44;	rity 34.0%; Pred. No. 1.7e-52; nservative 34; Mismatches 386; Ind	Best Loca Matches
	agen alpha l(IV) chain 23.6%; Score 984; DB	C;Supertamily Query Match
	ce: scrain no/xv	C;Genetics: A;Gene: Rv057
AL123456; NIC	RESIDUES: 1-1306 CCOL>  RESIDUES: 1-1306 CCOL>  Cross-references: UNIFROT:053775; UNIFARC:UPI00000D4FC8; GB:AL021942; GB:	A;Residues: 1-1306 A;Cross-references:
משנ	on: A/0934  preliminary; nucleic acid sequence not shown; translation not shown  troe. Nur	A;Accession: A;Status: pre
3.G. mplete genome	~ a a	A; Authors: Sqares, A; Title: Decipherin A; Reference number:
; Holroyd, S.	.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares,	; Connor, R.; Rajandream, M

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1130	765	1075	723	1015	696	955
1130 GTGGTGGAGGAGGTGS 1145	GVGPGAGAGSGAGAGS 780	1075 SPTGNNGSQGAGGDGGAGGAGGTGGTGGDGGRGAHGTLFSSLAGTGGTGGNGGTG- 1129	GAGSGAGAGSGVGVPGVGVGVGVGVGGVGGVGPGVGPGVGP 764	1015 GGAGGTGGTGLRGTTGATGATGTFDAGADGHGGNGGTGGVGGTGGAGGGGGGGGGGGKAL 1074	PGVGPGVGPGV 722	

Search completed: May 10, 2006, 10:44:30 Job time: 51 secs

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Result
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Maximum Match 100%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Scoring table:
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          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Q8VIY9 MYCTU
Q7TWBE MYCBO
Q6MWW7 MYCTU
Q7TWCB MYCBO
Q8WWG MYCBO
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O44359_NEPCL
Q8C9L8_MOUSE
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Q9BIT2_PLETR
Q9NHW4_NEPCL
Q9BIT7_9ARAC
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MEDLINE=89094868; PubMed=3210244; Mita K., Ichimura S., Zama M., James T.C.; Wita K., Ichimura S., Zama M., James T.C.; Septential Codon usage pattern and its implications structure of silk fibroin mRNA."; J. Mol. Biol. 203:917-925(1988).

on the secondary

PARTIAL NUCLEOTIDE SEQUENCE STRAIN=Kinshu X Showa;

"Highly repetitive structure and gene.";
J. Mol. Evol. 38:583-592(1994).

its

organization of the silk fibroin

DOI=10.1007/BF00175878;

NUCLEOTIDE SEQUENCE OF 5179-5263, STRAIN=J-139;

AND DISULFIDE BONDS

PARTIAL NUCLEOTIDE SEQUENCE.
MEDLINE=94365842; PubMed=7916056;
MITAK., IChimura S., James T.C.;

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Xia Q., Zhou Z., Lu C., Cheng D., Dai F., Li B., Zhao P., Zha X., Cheng T., Chai C., Pan G., Xu J., Liu C., Lin Y., Qian J., Hou Y., Wu Z., Li G., Pan M., Li C., Shen Y., Lan X., Yuan L., Li T., Xu H., Yang G., Wan Y., Zhu Y., Yu M., Shen W., Wu D., Xiang Z., Yu J., Wang J., Li R., Shi J., Li H., Li G., Su J., Wang X., Li G., Zhang Z., Wu Q., Li J., Zhang Q., Wei N., Xu J., Sun H., Dong L., Liu D., Zhao S., Zhao X., Meng Q., Lan F., Huang X., Li Y., Fang L., Li C., Li D., Sun Y., Zhang Z., Yang Z., Huang Y., Xi Y., Qi Q., He D., Huang H., Zhang X., Wang Z., Li W., Cao Y., Yu Y., Yu H., Li J., Ye J., Chen H., Zhou Y., Liu B., Wang J., Ye J., Ji H., Li S., Wang J., Wang J., Zhang Y., Zhang Y., Zheng H., Mao B., Wang W., Ye C., Li S., Wang J., Wang G.K.-S., Yang H., Mao B., Wang W., Ye C., Li S., Wang J., Wang G.K.-S., Yang H.,
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EMBL; V00094; CAA23432.1; -; Genomic_DNA.
EMBL; V00097; CAA23433.1; -; Genomic_DNA.
EMBL; S74439; AAB31861.1; -; mENA.
EMBL; X13869; CAA32076.1; -; mENA.
EMBL; M35378; AAA27839.1; -; mENA.
EMBL; M35378; AAA27839.1; -; mENA.
EMBL; M35378; BAA33147.1; -; Genomic_DNA.
EMBL; CK538369; -; NOT_ANNOTATED_CDS; MENA.
EMBL; CK538369; -; NOT_ANNOTATED_CDS; Genomic_DNA.
EMBL; AADK01000575; -; NOT_ANNOTATED_CDS; Genomic_DNA.
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19:6:
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Takagi T., Mizuno S.;
"Determination of the site of disulfide linkage
light chains of silk fibroin produced by Bombyx
Blochim. Biophys. Acta 1432:92-103(1999).
                                                         Repeat; Signal;
SIGNAL 1
CHAIN 22
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Tanaka K., Kajiyama N., Ishikura K., Waga S., Kikuchi A., Ohtomo K.,
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TISSUE
                                                                                                                                                                                                                                                                                                                                                                             s Swiss-Prot entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the El European Bioinformatics Institute. There are no rest
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE SPECIFICITY: Produced exclusively in the posterior (PSG) section of silk glands, which are essentially modified salivary glands.

Composed of antiparallel beta sheets. The strands of the beta sheets run parallel to the fiber axis. Long stretches of silk fibroin are composed of microcrystalline arrays of (-Gly-Ser-Gly-Ala-Gly-Ala-)n interrupted by regions containing bulkier residues. The fiber is composed of microcrystalline arrays alternating with amorphous regions.

PTW: The interchain disulfide bridge is essential for the intracellular transport and secretion of fibroin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and chemically inert fiber. SUBUNIT: Silk fibroin elementary unit consists in a disulfic SUBUNIT: Silk fibroin elementary unit consists in a disulfic linked heavy and light chain and a p25 glycoprocein in molar ratios of 6:6:1. This results in a complex of approximately
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Fibroin heavy chain.
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RESULT

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ID Q5Q294;
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GQ294;
05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Major ampullate spidroin (Fragment).
Agelenopsis aperta (Funnel-web spider).
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida;
Araneomorphae; Entelegynae; Agelenidae; Agelenopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biomacromolecules 0:0-0(2004).
EMBL; AY566305; AAT08436.1; -; mRNA.
NON_TER 1 1
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Tian M., Liu C., Lewis R.;
"Analysis of Major Ampullate Silk
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 GAGSGSGAAA-
                                         GT-GAGA
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                                        -GAGAGSGAGAGSGAGAGPGYGYAAYYAANYGYGPGVGAGSGAGAGSGSGSGA
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                                                                               -GAGAGSGAGAGSGAGAGGSGSGAGAGRGAGLGGLGAGLGSGV
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-GSGAGLGRGAGTGAGAGSGAGSGAGSGAGAGSGS
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RESULT 3
Q6PY84 KUKHI P
Q6PY84 KUKHI P
AC Q6PY84;
DT 05-JUL-2004 (T
DT 05-JUL-2004 (T
DT 05-JUL-2004 (T
DT 05-JUL-2004 (T
DE Major ampullat
OS KUKulcania hib
OC Eukaryota; Met
OC Arancomorphae;
OX NCBI_TaxID=268
RN [1]
RP NUCLEOTIDE SEQ
RA Tian M., Liu C
RA TIAN M
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Best Local Similarity
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Major ampullate spidroin 1 (Fragment).
Kukulcania hibernalis (Southern house spider).
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachmida;
Araneomorphae; Haplogynae; Filistatidae; Kukulcania.
MCBI TaxID=268415;
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"Analysis of Major Ampullate
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     AGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVFGKGVPGVGPGVGPGVGPGVGPGAG
                                                 RRGPGGARSRGGAGAGAGAGSGVGGYGSG-SGAGA-----GAGAGAGAGGEGGFGEGQG
                                                                                                                                                       AG---
                                                                                                                                                                                                      PGVGVPGVGVPGKG------VPGVGPGVGPG-VGPGVGPGAGAGSGAGAGSG
                                                                                                                                                                                                                                                                                  PGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGSGAGAGSGAGAGSGVGV
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760 AA;
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ilarity 46.3%;
Conservative 3
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                                                                                                   ----AGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGVGPGVGPGAG
                                                                                                                                                     GRGRGGRGGEAFSASSASSAVVFESAGPGEEAGSSGGGASAAASAAAAAGAGSG
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; 59989 MW; D98439784D9A2A8D
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Pred. No. 1.5e-84;
37; Mismatches 224; Indels 204;
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Q9BIUB_ARGTR
Q9BIUB_ARGTR PRELIMINARY; PRT; 1002 AA.
AC Q9BIUB;
DT 01-JUN-2001 (TrEMBLrel. 17, Created)
DT 01-JUN-2001 (TrEMBLrel. 24, Last sequence update)
DT 01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
DT 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
DE Plagelliform silk protein (Fragment).

Regiope trifasciata (Banded garden spider).
OC Araneomorphae; Entelegynae; Araneoidea; Araneidae; A.
OC Araneomorphae; Entelegynae; Araneoidea; Araneidae; A.
OX NCBI_TaxID=156845;
RN NUCLEOTIDE SEQUENCE.
RX MEDLINE=21179804; PubMed=11283372; DOI=10.1126/science ditersity, conservation, and convergence of fibroin sequences.";
Science 291:2603-2605(2001).
RP NUCLEOTIDE SEQUENCE.
RA Gatesy J.E., Hayashi C.Y.;
RA Go; GO:0006817; P:phosphate transport; IEA.
DR GO; GO:0006817; P:phosphate transport; IEA.
DR GO; GO:0006817; P:phosphate transport; IEA.
FT NON_TER 1002 1002
SEQUENCE 1002 AA; 75889 MW; 432D73E1EDF6A88A CRC
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Matches 392
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Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachmida; Ara
Araneomorphae; Entelegynae; Araneoidea; Araneidae; Argiope.
NCBI_TaxID=156845;
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   GPGVGPGAGAGSGAGAGSGAGAGSGVGVPGVPGVGVPGV----
                                     PGGPGGPGGPGGVGPGGAGGYGPGGAGGVGPAGTGGFGPGGAGGFGPGGAGGFGPGG
                                                                                                          GAGPGGAGGEGPVTVDVDVTVGPEGVG-GGPGGAGPGGAGFGPGGAGFGPGGAPGAPGG
                                                                                                                                           GAGSGAGAGSG-----
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YGAGAGAGFGSGAGAGAGAGAGA---
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                                                                                                                                                                             Conservative
                                                                    - GAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVGPGVGPGV
                                                                                                                                                                                          35.0%;
                                                                                                                                                                           Score 1459.5;
Pred. No. 3.6e.
3; Mismatches
                                                                                                                                       432D73E1EDF6A88A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; DOI=10.1126/science.1057561; , Woods J., Lewis R.; and convergence of spider silk
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                                                                                                                                                                                             5; DB 2;
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RESULT 5

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ID 017434;

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DT 01-JAN-1998 (T
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DT 01-JAN-1998 (T
DE Minor ampullate
OS Nephila clavip
OC Eukaryota; Met
OC Araneomorphae;
OC Araneomorphae;
OX NCBI_TaxID=691
RN [1]
RP NUCLEOTIDE SEQ
RY MEDLINE=982004
RA COlgin M.A., L
RT "Spider minor
RI "Spider minor
RI Protein Sci. 7
DR EMBL; AF027735
FT NON_TER
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01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Minor ampullate silk protein MiSp1 (Fragment).
Mephila clavipes (Orb spider).
Eukaryota, Metazoa, Arthropoda; Chelicerata, Arachnida;
Araneomorphae; Entelegynae; Araneoidea; Tetragnathidae;
                                Colgin M.A., Lewis R.V.;
"Spider minor ampullate silk proteins contain new and highly conserved non-silk-like 'spacer regions protein Sci. 7:667-672(1998).
                                                                                                                                                     NCBI_TaxID=6915;
 EMBL; AF027735;
NON_TER 1
                                                                                                      MEDLINE=98200471; PubMed=9541398;
                                                                                                                         NUCLEOTIDE SEQUENCE
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tive 46; Mismatches 286
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RA Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
RA Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
RA Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
RA Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
RA Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
RA Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
RA Micaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
RA Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
RA Nicaud J.-M., Nikolski M., Oztas S., Traub M.-L., Suleau A.,
RA Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
RA Zeniou-Meyer M., Zivanovic Y., Bolotin-Fukuhara M., Thierry A.,
RA Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
Wincker P., Souciet J.-L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local Similarity
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Similarities with tr|Q08294 Saccharomyces cerevisiae
OrderedLocusNames=CAGLOJ01771g;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Genome evolution in yeasts.";
Nature 430:35-44(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Candida glabrata (Yeast) (Torulopsis glabrata).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; mitosporic Saccharomycetales; Candida.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=ATCC 2001 / CBS 138;
PubMed=15229592; DOI=10.1038/nature02579;
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AGSGAGA--GSGAGAGSGVGVPGVGVPGVGVPGVGPGVGPGVGPGVGPGVGPGAGAGSG
                                                                                                                                                                     GAGAGSGAGAGSGVGVPGVGVPGVGVPGXGVPGVGPGVGPGVGPGVGPGAGAGSGAGAGS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 1397.5; DB 2; Pred. No. 2.6e-76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genomic_DNA.
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                                           ----STPGEG
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                                           SGSGSGSGSGSTPGEGSGSGSG
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483 180

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120 375 60

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RESULT 7
QNHW2 9A
ID Q9NH
AC Q9NH
AC Q9NH
DT 01-
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Best Local Similarity
Matches 412; Conserv
                                                                                                                                                                                                                                                                                                                                                                                        gene.";

Science 287:1477-1479(2000).

EMBL; AF218633; AAF36091.1; -; Genomic_!
GO; GO:0005737; C:cytoplasm; IEA.
GO; GO:0006817; P:phosphate transport;
InterPro; IPR008160; Collagen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vi-UCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Flagelliform silk protein (Fragment).
Name=Flag;
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Q9NHW2;
01-OCT-2000 (
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Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Araneae;
Araneomorphae; Entelegynae; Araneoidea; Tetragnathidae; Nephila.
NCBI_TaxID=115969;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE.
MEDLINE=20156766; PubMed=10688794;
Hayashi C.Y., Lewis R.V.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Molecular architecture and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9ARAC
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     vgp-gvgp-gvgp-gvgpg
                                                                           GPGGSGPGGAGSGGVGPGGYGPGGAGPG-GAGPGGAGPSGAGPGGAGTGGAGTGG
                                                                                                      GPG----AGAGSG------AGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPG
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1884 AA;
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ilarity 42.3%; Pred. No. 2.6e
Conservative 16; Mismatches
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                               ----AGAG-SGAGAGSGAGAGSGAGA
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ELN CHICK
ID ELN CHICK
AC P07916;
DT 01-AUG-1988 (
DT 01-AUG-1988 (
DT 10-MAY-2005 (
DE Elastin precu
GN NamesELN;
OS Gallus gallus
OC Eukaryota; Me
OC Archosauria;
OC Gallus.
TaxID=90
RN [1]
RP NUCLEOTIDE SE
RX MEDLINE=87242
RA Bressan G.M.;
RT "Repeating st
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NUCLEOTIDE SEQUENCE.
MEDLINE=87242320; PubMed=3593675;
Bressan G.M., Argos P., Stanley K.K.;
"Repeating structure of chick tropoelastin
                                                                                                                                                                                                        01-AUG-1988 (Rel. 08, Created)
01-AUG-1988 (Rel. 08, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Elastin precursor (Tropoelastin) (Fragment).
                                                                                                                                                 Gallus gallus (Chicken).
Eukaryota; Metazoa; Chor
                                                                                           NCBI_TaxID=9031;
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InterPro; IPR009765; Pericardin rpt.
InterPro; IPR003979; troppelastin.
Pfam; PF01391; Collagen; 1.
Pfam; PF07054; Pericardin rpt; 2.
PRINTS; PR01500; TROPOELASTIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; M18633; AAA48761.1; -;
EMBL; M21880; AAA49082.1; -;
EMBL; M15889; AAA49108.1; -;
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Biochemistry 26:1497-1503(1987).
[2]
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SUBCELLULAR LOCATION: Extracellular matrix of elastic fibers.
ALTERNATIVE PRODUCTS:
Event=Alternative splicing; Named isoforms=2;
Comment=Additional isoforms seem to exist;
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M: The cross?
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RA George R. A., Lewis S. E., Richards S., Ashburner M., Henderson S. N., Ra Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Ra Brandon R.C., Rogers Y. H.C., Blazej R.G., Champe M., Pfeiffer B.D., RA Brandon R.C., Rogers Y. H.C., Blazej R.G., Champe M., Pfeiffer B.D., RA Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P., Brottier R., Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., RA Dodson K., Dough L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., RA Dodson K., Dough L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., RA Horstin M., Evangelista C.C., Ferriac C., Ferriac S., Fleischmann W., RA Fosler C., Gabriellan A.E., Garg N.S., Gelbart W.M., Glasser K., Andrew D.A., Heiman T.J., Hernandez J.R., Houck J., RA Horstin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C., Karberis N., Harris M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A., RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z., Lang Y., Lin X., RA Lasko P., Levitsky A.A., Li J.H., Li Z., Liang Y., Lin X., RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M., RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M., RA Rainert K.M., Remington K., Saunders R.D.C., Scheeler F., Shen H., RA Spier E., Spradling A.C., Stapleton M., Strong R., Wang A.H., Wang X., RA Milliams S.M., Woodsey T., Worley K.C., Wang A.H., Wang X., RA Milliams S.M., Woodsey T., Worley K.C., Wang A.H., Wang X., Ra Alli M., Kalush F., Karpen G., Shan M., Strong R., Yao Q.A., Shith H.O., Shan M., Shan Q., Zhao Q.,
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Drosophila melanogaster (Fruit fly).
Bukaryota, Metazoa, Arthropoda; Hexapoda; Insecta; Pterygota;
Bueoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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GO; GO:0007507; P:heart development; IMP.
GO; GO:0002009; P:morphogenesis of an epithelium;
InterPro; IPR009765; Pericardin rpt.
Pfam; PF07054; Pericardin_rpt; 34.
Pfam; PF07054; Pericardin_rpt; 34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Celniker S.E., Wheeler D.A., Kronmiller B., Carlson J.W., Halpern A., Patel S., Adams M., Champe M., Dugan S.P., Frise E., Hodgson A., George R.A., Hoskins R.A., Laverty T., Muzny D.M., Nelson C.R., Pacleb J.M., Park S., Pfeiffer B.D., Richards S., Sodergren E.J., Svirskas R., Tabor P.E., Wan K., Stapleton M., Sutton G.G., Venter C., Weinstock G., Scherer S.E., Myers E.W., Gibbs R.A., Rubin G.M.; "Finishing a whole-genome shotgun: release 3 of the Drosophila melanogaster euchromatic genome sequence."; Genome Biol. 3:RESEARCH0079-RESEARCH0079(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=22426069; PubMed=12537572;
Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E.,
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B.J.
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
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Celniker S., Carlson J., Wan K., Pfeiffer
Hoskins R., Stapleton M., Pacleb J., Park
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Kaminker J.S., Bergman C.
Patel S., Frise E., Wheel
Ashburner M., Celniker S.
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Submitted (MAR-2000) to t
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genome sequence of Dros
Ice 287:2185-2195(2000).
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                                                                                              GYGSQLGVGAQAGAGQPGYGDQPGVGTQPGVGQPGYGSQPGTGAGQPGYGSQPGTG
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PubMed=12537573;

PubMed=12537573;

Lewis S.E.,
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EMBL/GenBank/DDBJ
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RESULT 10
Q9B1T2 PLETR PRELIMINARY; PRT; 912 AA.
ID Q9B1T2;
AC Q9B1T2;
DT 01-UN-2001 (TrEMBLrel. 17, Created)
DT 01-UN-2001 (TrEMBLrel. 17, Last sequence update)
DT 01-CCT-2002 (TrEMBLrel. 22, Last annotation update)
DE Fibroin 1 (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
MEDLINE=21179804; PubMed=11283372;
Gatesy J., Hayashi C., Motriuk D.,
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Araneomorphae; Haplogynae; Plectreuridae; Plectreurys.
NCBI_TaxID=33319;
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Science 291:2603-2605(2001)
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      GAGGAGAGFGSGLGLGYGVGLSSAQAQAQSAAAARAQADAQAQAYAAAQAQAQAQAQAQAQ
                              GVPGVGPGVGPGVGPG---------
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RESULT 11

Q9NHW4;

ID Q9NHW4;

AC Q9NHW4;

DT 01-OCT-2

DT 01-OM-2

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DE Flagelli

GN Name=Fla

OS Nephila

OC Eukaryot

OC Araneomo

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RR NUCLEOTI

RX MEDLINE=

RA Hayashi

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RL Science

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Science 287:1477-1479(2000).
Science 287:1477-1479(2000).
EMBL; AF218621; AAF36090.1; -; Genomic_DNA.
GO; GO:0005737; C:cytoplasm; IEA.
R GO; GO:0006817; P:phosphate transport; IEA.
R InterPro; IPR000160; Collagen.
InterPro; IPR000209; Pept S8 S53.
PR PROSITE; PS00138; SUBTILASE_SER; UNKNOWN_1.
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Best Local Similarity
Matches 418; Conserv
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Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Araneae;
Araneomorphae; Entelegynae; Araneoidea; Tetragnathidae; Nephila.
NCBI_TaxID=6915;
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01-OCT-2000 (TYEMBLrel. 15, Last sequence update)
01-WAR-2004 (TYEMBLrel. 26, Last annotation update)
Flagelliform silk protein (Fragment).
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Q9NHW4;
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Hayashi C.Y., Lewis R.V.;
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                             GGAGGAGGSGGTTIVEDLDITIDGADGPITISEELTIGGAGAGGSGPGGAGPGNVGPGRS
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Pred. No. 1.6e-70;
B; Mismatches 342;
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RESULT 12
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ID Q9BIT7;
O1-JUN-2001 (TrEMBLrel. 17, Created)
DT 01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
DT 01-JUN-2001 (TrEMBLrel. 24, Last annotation update)
DT 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
DT 01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
DE Major ampullate spidroin 2-like protein (Fragment).
OS Nephila inaurate madagascariensis.
OS Nephila inaurate madagascariensis.
OC Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Araneae;
OC Araneomorphae; Entelegynae; Araneoidea; Tetragnathidae; Nephila.
OX NCBI TaxID=115969;
RN NUCLEOTIDE SEQUENCE.
RX MEDLINE=21179804; PhDMed=11283372; DOI=10.1126/science.1057561;
RA Gatesy J., Hayashi C., Motriuk D., Woods J., Lewis R.;
RT fibroin sequences.";
Csience 291:2603-2605(2001).
RN 121
RP NUCLEOTIDE SEQUENCE.
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Submitted (FBB-2001) to the EMBL,

EMBL; AF350276; AAK30605.1; -; G

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SEQUENCE 1953 AA; 159383 MW;
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                                                  QGPGAAAAAAAGSGPGGYGPGQQGPGRSGAAAAAAAAAGRGPGGYGPGQQGPGGPGAAAAAA 1451
                                                                                                               SGPGGYGPGQQGPGGPGAAAAAAAAGRGPGGYGQGQQGPGGPGAAAAAAAAGPG---GYGPGQ
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AA; 159383 MW;
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                                                                                                                                                                                                      -GKGVPGVGPG-------VGP-GVGPG-VGPGAGAGSGAG
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Pred. No. 3e-70;
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Best Local Similarity
Matches 406; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                    Development 129:3241-3253(2002).

EMBL; AF203342; AAF11280.1; -; mRNA.

FlyBase; FBgn0028573; pcc.

GO; GO:0005578; C:extracellular matrix (sensu Me: GO; GO:0007507; P:heart development; IMP.

InterPro; IPR009765; Pericardin rpt.

Pfam; PF07054; Pericardin rpt; 33.

SEQUENCE 1729 AA; 164661 MW; DA9B1B7FABSCDED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=Oregon R;
MEDLINE=22065272; PubMed=12070098;
MEDLINE=22065272; PubMed=12070098;
Chartier A., Zaffran S., Astier M., Semeriva M., Gratecos D.;
"Pericardin, a Drosophila type IV collagen-like protein is involute morphogenesis and maintenance of the heart epithelium during dorsal ectoderm closure.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Name=prc; ORFNames=CG5700;
Drosophila melanogaster (Fruit fly).
Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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IT DROME
OBUGAT, DROME PRELIMINARY;
OBUGAT,
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                                                  GSQPGIGGQTGAGQPGYGTQPGIGGQTGAGQPGYGTQPGIGAQTGAGQPGYGTQPGIGGQ
                                                                                GSGAGAGSGAGAGS-----GAGAGSGVGVPGVGV-PGVGV-PGKGVPGVG--PGVG--
                                                                                                               QTGAGQPGYGTQPGIGGQTGAGQPGYGTQPGIGGQTGPGQPGYGTQPGIGGQTGPGQPGY
                                                                                                                                               GSGVGVPGVGV-PGV-GVPGKGVPGVG--PGVG------PGVGPGVGPGVGPG-AGA
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RESULT 14
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ID 044359,
AC 044359,
DT 01-JUN-1
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DT 01-JUN-1
CO Nephila
OS Nephila
OS Nephila
OC Eukaryott
OC Araneomo
OX NCBI_Tax
RN (1]
RP NUCLEOTI
RX MEDLINE-
RA Hayashi
RT "Evideno
RT elastici
RT "Buideno
RT elastici
RT EMBL, AF
                                                                                                                                                                                                                        O44359_NEPCL PRELIMINARY; PRT; 907 AA.

O44359;

O44359;

O1-JUN-1998 (TrEMBLrel. 06, Created)

O1-JUN-1998 (TrEMBLrel. 06, Last sequence update)

O1-JUN-2003 (TrEMBLrel. 24, Last annotation update)

Flagelliform silk protein (Fragment).

Name=Flag;
Nephila clavipes (Orb spider)

E Flagelliform silk protein (Fragment).

Nephila clavipes (Orb spider)

E Sukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Arachneomorphae; Entelegynae; Araneoidea; Tetragnathidae; Nephila Clavipes (Orb spider)
NUCLEOTIDE SEQUENCE.

MEDLINE-98153262; Pubmed=9480768; DOI=10.1006/jmbi.1997.1478;
Hayashi C.Y., Lewis R.V.;
"Evidence from flagelliform silk cDNA for the structural basi
"Estidence from flagelliform silk cDNA for the structural basi
elasticity and modular nature of spider silks.";
J. Mol. Biol. 275:773-784(1998)
EMBL; AF027973; AAC38847.1; -; mRNA.
GO; GO:0005737; C:cytoplasm; IEA.
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RC STRAINS—CSTBLINGUE=Thymus;
RC MEDLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;
RA Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,
RA Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,
RA Nikaido I., Osato N., Saito R., Suzuki H., Yyamanaka I., Kiyosawa H.,
RA Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,
RA Nikaido I., Osato N., Saito R., Schonbach C., Gojobori T.,
RA Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
RA Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
RA Balke J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA Balke J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA Balke J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA Gassterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,
RA Gassterland T., Gustincich S., Hirokawa N., Jacksson I.J., Jarvis E.D.,
RA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
RA Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
RA Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
RA Maglatt D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,
RA Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,
RA Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,
RA Ravasi T., Reed J.C., Reed D.J., Reid J., Ramachandran S.,
RA Petrovsky N., Pilai R., Pontius J.U., Qi D., Ramachandran S.,
RA Verardo R., Wagner L., Wahlestedt C., Wang Y., Watanabe Y., Wells C.,
RA Wilsana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M.,
RA Yuna Z., Zavolan M., Zhu Y., Zimmer A., Carninci P., Hayatsu N.,
RA Mining L.G., Wynshaw-Boris A., Yanagisawa M., Yang I., Yang L.,
RA Minashi T., Waki K., Kawai J., Akamura M., Sakazume N., Sakazume N., Sakazume N., Sakai K., Sakaki N., Saaski D., Shinagawa A.,
RA Yannishi A., Yashino M., Watsarton R., Lander R.S., Shinagawa I.,
RA Yannishi A., Yashino M., Watsarton R., Lander R.S., Shinagawa I.,
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
10-MAY-2005 (TrEMBLrel. 30, Last annotation update)
10-MAY-2005 (TrEMBLrel. 30, Last annotation update)
Mus musculus 3 days neonate thymus cDNA, RIKEN full-length enriched
library, clone:A630042I19 product:elastin, full insert sequence
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RA Strausberg R.L., Peingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Guellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Robask S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C STRAINCC57BL/6J; TISSUB=Thymus;

A Adachi J., Alzawa K., Akimura T., Arakawa T., Bono H., Carninci P.,

Fukuda S., Furuno M., Hanagaki T., Hara A., Hashizume W.,

Fukuda S., Furuno M., Hanagaki T., Hara A., Hashizume W.,

A Hayashida K., Hayatsu N., Hiramoto K., Hiraoka T., Hirozane T.,

A Hori F., Imotani K., Ishii Y., Itoh M., Kagawa I., Kasukawa T.,

A Katoh H., Kawai J., Kojima Y., Kondo S., Konno H., Kouda M., Koya S.,

A Kurihara C., Matsuyama T., Miyazaki A., Murata M., Nakamura M.,

A Nishi K., Nomura K., Numazaki R., Ohno M., Ohsato N., Okazaki Y.,

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A Saito R., Saitoh H., Sakai C., Sakai K., Sakazume N., Sano H.,

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Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
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MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
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"Analysis of the mouse transcriptome based 60,770 full-length cDNAs.";
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Submitted (APR-2003) to the EMBL; AK041860; BAC31084.1; EMBL; BC051649; AAH51649.1;

EMBL/GenBank/DDBJ databases

mRNA.

Strausberg

SORBRAR

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Query Match
Best Local Similarity
Matches 375; Conserv
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MGI; MGI:95317; Eln.
GO; GO:0007519; P:myogenesis; IMP.
GO; GO:0030833; P:regulation of actin filament polymerization; IMP.
GO; GO:0043149; P:stress fiber formation; IMP.
InterPro; IPR003979; troppelastin.
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Biocceleration Ltd.
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Silk-elas
Active ag
Silk-elas
SELP 47K
Protein p
SELP8K po
Expected
SELP8K po
Plasmid p
SELP8K po
Plasmid p
Silk-elas
Active ag
Silk/elas
SELP 47E-
Cotton bi
Cotton bi
Actimicro
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Silk-elas
Active ag
Silk/elas
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3182	3192	3250	3253	3253	3263	3263	3264	3284	3284	3296	3296	3296	3296	3296	3296	3296	3305	3305	3305	3305
76.3	76.5	77.9	78.0	78.0	78.2	78.2	78.2	78.7	78.7	79.0	79.0	79.0	79.0	79.0	79.0	79.0	79.2	79.2	79.2	79.2
988	768	1043	832	832	1016	1016	768	877	877	2257	2257	2257	2257	2257	2257	2257	1038	1038	1038	1038
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AAR80253	ABP53481	ADU81676	ABP53473	AAR80252	AEB57388	ADU81677	ABP53466	AAW49724	AAR80335	ADE44977	ABG69272	AAY78282	AAW53523	AAW26347	AAR41012	AAP82961	AEB57386	ADU81671	ADU98740	ADR70478
Aar80253	Abp53481		Abp53473	Aar80252	Aeb57388	Adu81677	Abp53466	Aaw49724	Aar80335	Ade44977	Abg69272	Aay78282	Aaw53523	Aaw26347	Aar41012	Aap82961	Aeb57386	Adu81671	Adu98740	Adr70478
Polymer S	Protein p	Cotton bi	Protein p	Polymer S	SELP47K-P	Radioprot	SELP8 rel	Protein p	Protein p	Recombina	Silk/Elas		Amino aci	SELP3 syn	SELP3 mul	SEPL3 pro	SELP 67K	Silk/elas	Active ag	Silk-elas

ALIGNMENTS

ADK51951 standard; protein; 780 AA

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WPI; 2004-023263/02.
                                                                                                                  04-DEC-2003
                                                                                                                                                 Repeat
                                                                                                                                                         06-MAY-2004
                                                                                                                                                                 ADK51951;
                                                                               Kumar M;
                                                                                                  20-MAY-2002; 2002US-0381913P
                                                                                                          20-MAY-2003; 2003WO-US015757.
                                                                                                                          WO2003099465-A1
                                                                                                                                  Unidentified
                                                                                                                                         repeat protein.
                                                                                       (DOWO ) DOW CORNING CORP.
                                                                                                                                                 protein polymer repeat sequence,
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The present invention relates to a method for forming an inorganic structure. The method comprises providing substrate with a repeat protein polymer, and exposing the substrate to a precursor comprising inorganic species. The repeat protein polymer catalyzes the formation of an inorganic structure on the substrate. The repeat portion of the repeat protein polymer may be ANKS1951-ADKS1957, which may have a head sequence at the N-terminal end (ADKS1958) and a tail sequence at the C-terminal Forming inorganic structure, for use as e.g. electronic photonic and national composite materials, comprises providing substrate with repeat protein polymer, and exposing substrate to precursor containing inorganic Claim 12; Page species. 8; 27pp; English.

Sequence 780 AA;

end. (ADK51959-ADK51966).

Aeb57381 Adu81678 Adu81673 Adu81674

Adu81675 Adr70479 Adu98741

Length

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RESULT 2
AD70468
ID 70468
ID 70470
XX ADR7
XX ADR7
XX YO-D
XX O2-D
XX Silk
XX Perf
KW hair
KW styl
KW anti
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Best Local Similarity
Matches 780; Conserv
 personal care composition; repeat sequence protein polymer; hair care composition; shampoo; conditioner; anti-dandruff treatment; styling aid; styling conditioner; hair repair treatment; chemical hair treatment; hair dye; skin care composition; body wash; antimicrobial cleanser; skin protectant cream; body lotion; facial cr moisturising cream; facial cleansing emulsion;
                                                                                                          Silk-elastin
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                                                                                                          polymer SELP47K
                                                                                                                                      (first entry)
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anti-acne treatment; facial toner; exfoliating cream; facial mask; after shave balm; sunscreen; anti-fungal treatment; skin protectant; antiperspirant; cosmetic composition; makeup; eye gel; high-melting point lipstick; lipstick; lip gloss; lip balm; mascara; eyeliner; pressed powder formulation; foundation; nail care composition; nail enamel; cuticle treatment; nail polish; nail treatment; polish remover; oral care composition; toothpaste; mouth rinse; breath freshener; teeth whitening treatment; over-the-counter pharmaceutical composition; transparent film formation; hydrogel formation; water-solubility; luster; softness; moisture retainment; tensile property; viscoelastic behaviour; glass transition temperature; cloud temperature; glass transition temperature; silk; elastin; repeating sequence unit.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2004180027-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-MAR-2003; 2003US-0454077P.
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Personal care composition useful as cosmetic, hair care or skin care product, comprises repeat sequence protein polymer and compounds such carriers, excipients, liposomes, active ingredients, or emollients. 28

WPI; 2004-675584/66

Kumar M,

(CUEV/) CUEVAS

ξ ₽

Example 1; SEQ ID NO 19; 50pp; English.

Crapeat sequence protein polymer with the balance of the composition accomprising ane or more compounds chosen from carriers, excipients, iliposomes, active ingredients, biological or botanical products, chamecrants, emollients, surfactants, thickening agents, silicone components, organic sunscreens, preservatives, neutralizing agents, conditioner, anti-dandruff treatment, styling aids, styling accomponents, organic sunscreens, preservatives, neutralizing agents, conditioner, anti-dandruff treatment, styling aids, styling accomponents, organic sunscreens, preservatives, neutralizing agents, conditioner, anti-dandruff treatment, styling aids, styling accomponents, organic sunscreens, preservatives, neutralizing agents, silicone components, styling aids, anti-acne composition such as anti-acne composition topically applied cover-the-counter formulation and foundation, and anti-perspirants, commetic composition such as nail care composition such as nail treatment, or polish remover care composition and foundation and treatment, or polish remover care and care composition such as nail treatment, or polish remover care composition is a mascara, water-based mascara, or composition such as mall treatment, or polish remover care and the styling aids, and the styling aids, and the styling aids at a star styling

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 Active agent
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ilarity 100.0%;
Conservative (
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system-related SELP47K protein SeqID19
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Pred. No. 4.4e-264;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention relates to a novel system for providing controlled release delivery of an active agent which comprises a repeat sequence protein polymer and at least one active agent. The invention may be useful in a hair care composition, skin care composition, nail care composition, cosmetic composition, oral care composition or over-the-counter pharmaceutical composition. The system may be useful in shampoos, gels, mousses and other hair care products, rinse-off conditioners, skin care products such as moisturizers, toners and makeup and nail care products such as polishes and polish removers. The invention enables controlled release of active agents. The present sequence is that of an SELP47K protein which may be used in the system of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                System for providing controlled release delivery of active agent useful for incorporating active agents into personal care product compositions, comprises repeat sequence protein polymer and active agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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The invention relates to biomolecular conjugates comprising a repeat sequence protein polymer (i.e., an engineered protein) and at least one active agent, such as silicones, fragrances, dyes, tints, UV actives, sunscreens, lanolin, vitamins, bleaches, thickening agents, antimicrobials and biologically active polypeptides. The repeat sequence protein polymer preferably comprises a replyanting amino acid sequence unit derived from elastin, collagen, abductin, byssus, flagelliform silk, dragline silk, gluten high molecular weight subunit, titin, fibronecuin,
                                                                                                                                                                                                                                                                                                                                    Biomolecular conjugates useful in personal care products e.g. hair composition, commetic, oral care composition, comprising conjugation product of repeat sequence protein polymer and active agents.
                                                                                                                                                                                                                                                                                 Claim
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cc mucin, RNA polymerase II, resilin or mixtures thereof. The repeat commany be bound. In particular, they enhance the ability to utilize proteins cc may be bound. In particular, they enhance the ability to utilize proteins cc and peptides as active agents as they overcome many of the polymers provide a scraffolding to which active agents can peptide active agents, and allow the use of polypeptide active agents, and allow the use of polypeptide active agents in applications for which a total protein environment is desirable. The compositions also relates to personal care compositions comprising a composition also relates to personal care compositions comprising a correct protein polymer and a polypeptide active agent; biomaterials adapted for at least one predetermined desirable function which comprise one or more biomolecular conjugates comprising a silk elastin polymer as the repeat sequence protein polymer and a polypeptide active agent; and a method of producing such biomaterials. The biomolecular conjugates are useful in personal care compositions used for hair care, skin care, nail care, compositions, and compositions used for hair care, skin care, nail care, compositions, and dressings, and in biomaterials useful in genetics research tools or in search and/or identification tools. The present correct polymer comprising silk fibroin and elastin repeat sequence protein polymer comprising silk fibroin and elastin repeat sequences which is specifically claimed for use in biomolecular conjugates of the convertion.
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CC The invention relates to a personal care composition (I) comprising: (a) CC a repeat sequence protein polymer (RSPP) or its fragment, and a carrier CC or excipient, where repeat sequence protein polymer provides at least one CC commetic benefit; or (b) a RSPP or its fragment, and one or more CC commounds chosen from carriers, excipients, liposomes, biological or CC compounds chosen from carriers, excipients, liposomes, biological or CC fragment improves skin tone. (I) is useful for protecting skin from CC environmental damage, where the environmental damage comprises damage due to exposure to radiation (especially ultraviolet light) or damage due to care CC composition such as moisturizing body washes, body washes, antimicrobial CC cleansers, skin protective creams, body lotions, facial creams, CC composition skin protective creams, body lotions, facial creams, CC comprising creams, facial masks, aftershave balms, composition.
CC comprising creams, facial masks, aftershave balms, composition.
CC comprising tropically applied over-the counter compositions, anti-fungal treatments, skin protectants, anti-perspirants, or radioprotective such as sunscreen chosen from non-water-resistant sunscreens. (I) is useful as a makeup composition such as eye gels, high-melting point lipsticks, commistations chosen from loose powders, blushes, eye shadows, and cCC bronzing powders, foundation chosen from water-in-oil foundations, water-in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Personal care composition useful for protecting skin from environmental damage, has carrier and repeat sequence protein polymer capable of providing cosmetic benefit such as increasing skin fibroblast viability.
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12-MAR-2004; 2004US-00800179.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 780 AA;
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Pred. No. 4.4e-264;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein polymers comprising repeating units and enzyme-catalysed covalent bond formation useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pendent group; repeating unit; enzyme recognition site; sealant; fibrin; enzymatic cross-linking; biocompatible material; structural integrity; medical adhesive; wound closure; tissue repair; transglutaminase.
                                                                                                                                                                                                                                                                            Sequence 884 AA;
                                                                                                                                                                                                                                                                                                                                                                                                       Example 7; Page 65; 138pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                             material for wound closure and tissue repair.
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                                                                                                                                                                                                                             Score 3448; DB 2;
Pred. No. 6.9e-217;
0; Mismatches 26;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         sequences - capable as a biocompatible
                                                                                                                                                                                                                                                    Length 884;
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                                                                                                                                                                                                                             SELP8K; protein polymer; silk; fibroin; elastin; repeat sequence; crosslinking; adhesive; tissue sealant; biocompatible film.
                                                                Misc-difference
                                                                                   Peptide
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                                                                                                                                                             Peptide
                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                         SELP8K polymer.
                                                                                                                                                                                                                                                                            29-MAR-1997
                                                                                                                                                                                                                                                                                                                 AAW09213 standard;
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        /note= "Silk fibroin peptide repeat sequence"
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83
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"Elastin peptide repeat sequence"
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                                                                                                               "Block
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                                                                                                                                                                      "Silk fibroin
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                                                      functional group for crosslinking"
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                                    repeat sequence"
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Best Local :
                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                     This SELPBK polymer sequence (mol.wt. 69,772) has been constructed from monomers composed of repeat units derived from silk fibroin (AAW09206) and elastin (AAW09207) (claim 4). The monomer (AAW09212) has been amplified by multicopy cloning in Escherichia coli (using plasmid pPT0345) to produce a recombinant protein polymer backbone with Lys functional groups for subsequent polyfunctional chemical crosslinking. The crosslinked polymer product forms a strongly adherent tissue adhesive or sealant. The polymer may be used to seal defects in vessel walls, e.g. artery, vein, capillary, lung, dura or colon, to increase tissue mass, or to produce biocompatible films for in vivo use. The sealants have the biocompatiblity of fibrin glues, but set more quickly, and have greater strength. They are readily prepared by recombinant methods, are easy to administer, and are gradually resorbed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissue repair adhesive comprising polymer of structural protein repeat units - contg. hetero-atom functional gps. reactive with crosslinking agent, combines biocompatibility and high bonding strength.
                                                                                                                                                                                                                                                                                                                                                                 Sequence 884
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Similarity 90.6%;
52; Conservative
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GAGAGSGAGAGSGVGVPGVGVPGVGVPGVGVPGKGVPGVGVPGVGVPGVPGVPGVPGA
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Pred. No. 6.9e-217;
0; Mismatches 26;
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                                                                           Preparation of synthetic repetitive DNA - useful for construction large protein polymers having repeating units, used in structural material, e.g. synthetic silk.
                                                                                                                                                                                                                                 rorymer SELP8K; peptide repeat
high molecular weight polymer;
                                                                                                                                                                                                                                                                             10-AUG-1998
                                                                                                                                                                                                                                                                                             AAW53541;
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                                                             Example 7; Page 89; 127pp;
                                                                                                                           Ferrari
                                                                                                           WPI; 1998-193613/17.
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                                                                                                                           Crissman
                                                             English
                                                                                                                                                                                                                                                              of pPT0345 encoding
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This is the amino acid sequence of pPT0345 comprising the SELP8K protein, used in the method of invention, which involves the preparation of synthetic DNA sequence having repeating units from bout 3-15 codons and encoding a protein of at least about 30 kDa. The method is useful for the production of high molecular weight polymers (e.g. synthetic silk), either nucleic acids or peptides that are the expression products of the

nucleic acids

and particularly high molecular weight

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                                                                                                                                                                                                                                                   SELPBK polymer is a synthetic silk-like protein comprising multiple copies of a monomer (see AAW49726) consisting of repeating units of silk-like sequences (GAGAS) and elastin-like sequences (VPGVG) including a reactive lysine residue. It was expressed in Escherichia coli from plasmid ppr0345 as a 80 kDa protein. The SELPBK polymer was specifically designed to be functionalised with different reagents reactive to amines. Claimed recombinant protein polymers are capable of covalent crosslinking by enzymatic reaction to form products which set quickly and have good adhesive properties and high strength. The proteins can comprise a repetitive amino acid backbone of repetitive units having collagen, fibroin, elastin or keratin motifs and at least 2 enzyme recognition sequences comprising a glutamine or lysine capable of enzyme catalysed isopeptide formation. The products can be used as medical adhesives and sealants, in the closure of wounds and repair of damaged tissue. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New recombinant protein polymers - repetitive units for crosslinking land sealants, depots and matrices.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Protein polymer; SELP8K; silk-like protein; fibroin; elastin; adhesive; sealant; wound healing; transglutaminase; cross-linking.
                                                                                                                                                                                                                             Sequence 884 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 7; Col 43; 70pp; English
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                                                                                                                                                                                    82.6%;
90.6%;
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                                                                                                                                                                                    Score 3448;
Pred. No. 6.
                                                                                                                                                                        Mismatches

    containing naturally
by enzymes, useful as

                                                                                                                                                                                     .9e-217;
                                                                                                                                                                                                  DВ
                                                                                                                                                                                                  2
                                                                                                                                                                                                 Length 884;
                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           medical adhesives
                                                                                                                                                                        52;
                                                                                                                                                                      Gaps
  213
                              170
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RESULT 10
AAY51882
ID AAY51
XX AAY51
AC AAY51
DT 22-JI
DX Cros
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XX Synt
XX US61
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This invention describes a novel crosslinked protein composition (A) which, before crosslinking, the protein (I) is new and comprises at I 70 wt.% of repeating units GAGAGS (I) and GVGVP (2), and in at least repeating units an amino acid (aa) is substituted by Lys or Arg to provide a Lys/Arg equivalent weight of 1-20 kD. (I) contains at least
                                                                                                                                                                                Crosslinked protein composition, useful as tissue adhesive or comprises peptide repeating units that contain functional grouwith crosslinker.
                                                                                                                                           Example
                                                                                                                                                                                                                                                                                                                                                                                                                  05-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Crosslinked
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VGVPGVGYPGKGVPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PGKGVPGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein; fibrin glue; tissue adhesive;
                                                                                                                                                                                                                                                                                                                         Stedronsky
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                                                                                                                                           51-56;
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                                                                                                                                      45pp;
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                                                                                                                                      English.
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    aa having a functional group reactive with at least one of aldehyde, iso(thio) cyanate and activated carboxy. (I) have similar biocompatibilty to fibrin glues, but set more quickly and give a bond with greater shear strength. They are made from readily available natural sources, are easy to administer and are gradually resorbed. This sequence represents a crossslinking protein polymer unit designated SELPBK
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t Local 5-
752;
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                                                                                                                                        VPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGV
                                                                                                                                                                          GAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVG-PGVG-PGVG-PGVG
VGVPGVGVPGKGVPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGV
                                                                               GVGVPGKGVPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPG
                                                                                                 PGKGVPGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVP
                                                                                                                                 884 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                vGVPGKGVPGVGVPGVGVPGVGVPGAGAGSGAGAGSGAGAGSGAGAGSGVGVPG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              82.6%;
90.6%;
                                /PGKGVPGVGVPGVGVPGVGVPGAGAGSGAGAGSGAGAGSGAGAGSGV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3448; DB 3;
No. 6.9e-217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
               780
                                813
                                                                                                                                                 566
                                                                                                                                                                  573
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RESULT 11 ABG31412 ID ABG314

ABG31412 standard; protein; 884 AA.

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Best Local S
Matches 752
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                                                                                                                                                                                                                                                                                                                      The present invention relates to protein polymers having repetitive units from naturally occurring structural proteins such as fibroin, elastin, collagen and keratin. The polymers comprise a functional group which can be chemically crosslinked with appropriate crosslinkers. The protein polymer is produced by recombinant DNA technology. The protein polymer is useful for sealing or filling a defect in viable tissue, particularly for augmenting tissue mass. The protein polymer is useful in a variety of applications related to their physical, chemical and biological properties, and/or to bond together separated tissue to provide a stable, flexible or resorbable bond. The protein is particularly useful as a sealant or adhesive, in wound healing or as a burn dressing e.g. to stop or staunch the flow of fluid (e.g. blood) through ruptured vessels (e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Crosslinked protein composition used as sealant or adhesive for sealing or filling defect in viable tissue, as burn dressing, or in wound healing e.g. to staunch flow of fluid e.g. blood, through ruptured vessels.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stedronsky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-MAY-1995;
02-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US6423333-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SELP8K polymer encoded by plasmid pPT0345
                                                                                                                                                                                                                                                                                  Sequence 884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-NOV-2002
                                                                                                                                                                                                                                                                                                            arteries or veins).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (PROT-) PROTEIN POLYMER TECHNOLOGIES
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                           171
                                                      154
                                                                                                                                      59
                                                                                                            94
                                                                                                                                                                  34
                                                                                                                                                                                                                           752;
                                                                                                                                                                                  Similarity
                                                                    GAGAGSGAGAGSGVGVPGVGVPGVGVPGVGVPGKGVPGVGVPGVGVPGVPGVPGVPGA
                                                                                                                                                                    Col 17; 46pp; English.
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                                                                                                                                                                                                                             Conservative
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96US-00642246.
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                                                                                                                                                                                                                                                                                                            The present sequence represents SELP8K polymer
                                                                                                                                                                                                                                       82.6%;
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                                                                                                                                                                                                                          Score 3448; DB 5;
Pred. No. 6.9e-217;
0; Mismatches 26;
                                                                                                                                                                                                                                                    Length 884;
                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                           52;
                                                                                                                                                                                                                          Gaps
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273
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RESULT 12
ABW01628
ID ABW016
XX ABW01
XX ABW01
XX 12-FE
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PN US200
XX Unide
PN US200
XX 05-JU
XX 05-AP
PF 05-AP
PR 02-MA
PR 29-NO
XX 12-MA
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                                                                                                                                                               05-MAY-1995;
02-MAY-1996;
29-NOV-1999;
                                          WPI; 2003-829350/77
                                                                                                                                                                                                                                                 05-APR-2002; 2002US-00117931.
                                                                                                                                                                                                                                                                                          05-JUN-2003
                                                                                                                                                                                                                                                                                                                                US2003104589-A1
                                                                                                                                                                                                                                                                                                                                                                          Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                               coating; vulnerary.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Crosslinked protein; sealant; depot; drug; therapy; gel; film; thread;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Plasmid pPTO345 SELP8K polymer protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABW01628;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABW01628 standard;
                                                                                                                           (PROT-) PROTEIN POLYMER TECHNOLOGIES
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99US-00451206.
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Crosslinked protein composition useful as sealing a defect in tissue,

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Best Local
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                                                                                                                                                               PGKGVPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVP
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90.6%;
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Pred. No. 6.9e-217;
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The invention describes a personal care composition (I) comprising a CC repeat sequence protein polymer with the balance of the composition CC comprising one or more compounds chosen from carriers, excipients, CC liposomes, active ingredients, biological or botanical products, CC components, organic sunscreens, preservatives, neutralising agents, CC perfumes and pigmente. (I) is useful as a hair care composition such as CC gerfumes and pigmente, anti-dandruff treatment, styling aids, styling CC conditioner, hair repair or treatment, serum, lotion, cream, pomade, or CC chemical treatment such as hair days; skin care composition such as CC gertumes and pigments, body wash, antimicrobial cleanser, skin CC cleanser body wash, antimicrobial cleanser, skin cream, moisturising cream, facial CC cleansing body wash, body wash, antimicrobial cleanser, skin CC cleansing emulsion, surfactant-based facial cleanser, facial exfeliating CC gel, anti-acne treatment, facial troner, exfoliating cream, facial mask, CC after shave balm or sunscreen; skin care composition topically applied cover-the-counter drugs comprising anti-fungal treatments, anti-acne treatments, skin care composition topically applied cover-the-counter drugs comprising anti-fungal treatments, anti-acne treatments, skin corner, exfoliating composition topically applied cover-the-counter drugs comprising anti-fungal treatments, anti-acne composition chosen from eye gel, high-melting point
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KW personal care composition; repeat sequence protein polymer;

KW hair care composition; shampoo; conditioner; anti-dandruff treatment;

KW styling aid; styling conditioner; hair repair treatment;

KW chemical hair treatment; hair dye; skin care composition; body wash;

KW antimicrobial cleanser; skin protectant cream; body lotion; facial cream;

KW moisturising cream; facial cleanser; genulsion;

KW surfactant-based facial cleanser; exfoliating gel;

KW anti-acne treatment; facial toner; exfoliating cream; facial mask;

KW after shave balm; sunscreen; anti-tungal treatment; skin protectant;

KW antiperspirant; commetic composition; makeup; eye gel;

KW high-melring point lipstick; lipstick; lip gloss; lip balm; mascara;

KW eyeliner; pressed powder formulation; foundation; nail care composition;

KW moisture research care composition; toothpaste; mouth rinse;

KW breath freshener; teeth whitening treatment;

KW wore-the-counter pharmaceutical composition; transparent film formation;

KW wore-the-counter pharmaceutical composition;

KW moisture retainment; tensile property; viscoelastic behaviour;

KW decommedition temperature; cloud temperature; softness;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Personal care composition useful as cosmetic, hair care or skin car product, comprises repeat sequence protein polymer and compounds su carriers, excipients, liposomes, active ingredients, or emollients.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-675584/66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-MAR-2003; 2003US-0454077P
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(CUEV/) CUEVAS W A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-MAR-2004; 2004US-00800179
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                           This invention relates to a novel system for providing controlled release delivery of an active agent which comprises a repeat sequence protein polymer and at least one active agent. The invention may be useful in a hair care composition, skin care composition, nail care composition, cosmetic composition, oral care composition or over-the-counter pharmaceutical composition. The system may be useful in shampoos, gels, mousses and other hair care products, rinse-off conditioners, skin care products such as moisturizers, toners and makeup and nail care products such as spolishes and polish removers. The invention enables controlled release of active agents. The present sequence is that of a protein which is related to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            System for providing controlled release delivery of active agent useful for incorporating active agents into personal care product compositions
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(MAZE/) MAZEAUD I.
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Pred. No. 6.9e-217;
0; Mismatches 26;
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Vertebrata.
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                                                                                                                                                                                                                                engineering; protein stabilization; cosmetics; textile; filter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VGVPGVGVPGKGVPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGV
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                                                                                                                                                            CC sequence protein polymer (i.e., an engineered protein) and at least one cative agent, such as slicones, fragrances, dyes, tints, UV actives, CC sunscreens, lanolin, vitamins, bleaches, thickening agents, interpreted protein polymer preferably comprises a repeating amino acid sequence comprised from elastin, collagen, abductin, byssus, flagelliform silk, CC derived from elastin, collagen, abductin, byssus, flagelliform silk, CC dragline silk, gluten high molecular weight subunit, titin, fibronectin, CC leminin, gliadin, glue polypeptide, ice nucleating protein, keratin, CC dragline silk, gluten high molecular weight subunit, titin, fibronectin, CC mucin, RNA polymerase II, resilin or mixtures thereof. The repeat sequence protein polymers provide a scaffolding to which active agents agents agents agents. The personal care compositions associated with gently peptide as active agents as they overcome many of the CC hydrophobicity, solubility, and binding limitations associated with gently peptide active agents, and allow the use of polypeptide active agents. The cC in applications for which a total protein environment is desirable. The CC in applications as nebtod of making a biomolecular conjugate by recompositions; a method of making a biomolecular conjugate by recompositions and polypeptide active agent; biomaterials adapted for at least one predetermined desirable function which comprise one or more compositions such biomaterials active agent; and a method of producing such biomaterials. The biomolecular conjugate are useful in gersonal care campositions such as over-the counter pharmaceutical corporation and clastin repeat sequence protein polymer and a polypeptide active agent; and a method of producing such biomaterials. The biomolecular conjugates are also useful in generic sequence represents a repeat sequence. Note: The present sequence is given the sequence is given in the sequence listing, but is not further referred to in the
                                                                                Matches
                                                                                                                     Query Match
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(KUMA/)
                                                                                                                                                          Sequence 884 AA;
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                                                                          Score 3448; DB y; ...
Pred. No. 6.9e-217; ...
""matches 26;
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Search completed: May 10, 2006, 10:39:41 Job time: 195 secs

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Maximum Match 100%
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| Sequence 68, Application US/08397633A |
| Patent No. 5773577 |
| GENERAL INFORMATION: GENERAL INFORMATION: APPLICANT: Cappello, Joseph |
| TITLE OF INVENTION: PRODUCTS COMPRISING SUBSTRATESCAPABLE TITLE OF INVENTION: OF ENZYMATIC CROSS-LINKING NUMBER OF SEQUENCES: 105 |
| CORRESPONDENCE ADDRESS: ADDRESSE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT STREET: 4 Embarcadero Center, Suite 3400 |
| CITY: San Francisco STATE: California COUNTRY: USA ZIP: 94111-4187
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; TOPOLOGY: linear
; MOLECULE TYPE: protein
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INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 884 amino acids
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ATTORNEY/AGENT INFORMATION:
NAME: Rowland, Bertram I
REGISTRATION UNMBER: 20,015
REFERENCE/DOCKET NUMBER: A-58
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/397,633A
FILING DATE:
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Pred. No. 3.3e-248;
No. Mismatches 26;
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US-08-435-641-15
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APPLICANT: Stedronsky, Erwin R.
TITLE OF INVENTION: Tissue Adhesive Using
TITLE OF INVENTION: Crosslinking
                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 415-781-198
TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,641
FILING DATE: 05-MAY-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Downland Bartram 1
                                                                                                                                                                                                                                                                                              STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE: |
                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Rowland, Bertram I.
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A-61127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: rour CITY: San Francisco
                                                                                                                                                                                                                                       Local Similarity
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                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
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STREET: Four Embarcadero Center, Suite 3400
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Pred. No. 3.3e-248;
0; Mismatches 26;
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US-08-707-237A-96
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Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
FILING DATE: 03-SEP-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/175,155
FILING DATE: 29-DEC-1993
                                                                                                                                                                                                                                                                          APPLICANT: Ferrari, Franco A.
APPLICANT: Capello, Joseph
APPLICANT: Crissman, John W.
APPLICANT: Dorman, Mary A.
TITLE OF INVENTION: METHODS FOR PREPARING SYNTHETIC
TITLE OF INVENTION: REPETITIVE DNA
                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: ISM PC COMPATIBLE
COMPUTER: PM PC POS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                            NUMBER OF SEQUENCES: 108
CORRESPONDENCE ADDRESS:
CORRESPONDENCE Flehr, Hohbach, Test,
STREET: Four Embarcadero Center,
CITY: San Francisco
STATE: California
                                                                                                                                                                               STATE: C
                                                                     APPLICATION NUMBER: US/08/707,237A
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Suite 3400
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FILING DATE: 29-OCT-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 06/927,258
FILING DATE: 04-NOV-1986
ATTORNEY/AGENT INFORMATION:
NAME: Trecarctin, Richard F.
REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-55186-10/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEPAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 96:
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APPLICATION NUMBER: US
FILING DATE: 22-APR-1S
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 06-NOV-1S
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Similarity 90.6%;
52; Conservative
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amino acid
VPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGV 566
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Pred. No. 3.3e-248;
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                                                                                                                             Query Match
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APPLICANT: STEDRONSKY, Erwin R.
APPLICANT: CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive
TITLE OF INVENTION: Crosslinking
                                                                                                                                                                                                                                   TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: A6
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION NUMBER: US/08/642,246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST,
STREET: Four Embarcadero Center,
                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: CA
COUNTRY: US
ZIP: 94111
                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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CITY: San Francisco
                                                                                                                 Local Similarity
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                                                                                                                                                                                                                      LENGTH: 884 amino acids
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                                                                                                     Score 3448; DB 2;
Pred. No. 3.3e-248;
0; Mismatches 26;
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Suite 200
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RESULT 6
US-09-451-206-15
; Sequence 15, Application US/09451206
; Patent No. 6423333
; GENERAL INFORMATION:
                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST,
STREET: Four Embarcadero Center,
CITY: San Francisco
STATE: CA
COUNTRY: US
ZIP: 94111
        COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                           APPLICANT: STEDRONSKY, Erwin R.
CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive
  CURRENT APPLICATION DATA:
                                                                                                                         NUMBER OF SEQUENCES: 35
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Suite 200
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TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ
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nes 752; Conserv
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PRIOR APPLICATION DATA:
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                                                                  GAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGVPGKGVPGVG-PGVG-PGVG-PGVG
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REFERENCE/DOCKET NUMBER: A61127-1/BIR
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Pred. No. 3.3e-248;
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PCT-US96-06229-15
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GENERAL INFORMATION:
APPLICANT: CIEDRONSKY, Erwin R.
APPLICANT: CIEDRONSKY, Joseph
TITLE OF INVENTION: Tissue Adhesive Using Synthetic
TITLE OF INVENTION: Crosslinking
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: FOUR Embarcadero Center, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A611
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/06229
                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 884 amino acids
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Conservative
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Pred. No. 3.3e-248;
0; Mismatches 26;
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US-08-175-155-47
; Sequence 47, Ap
; Patent No. 5641
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          APPLICATION NUMBER: US/08/17
FILING DATE: 29-DEC-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Rowland, Bertram I.
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A-5
                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                           APPLICANT: Ferrari, Franco A.
APPLICANT: Cappello, Joseph
APPLICANT: Crissman, John W.
APPLICANT: Dorman, Mary A.
ATITLE OF INVENTION: Methods for Preparing
TITLE OF INVENTION: Repetitive DNA
NUMBER OF SEQUENCES: 69
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CITY: San Francisco
STATE: CA
COUNTRY: US
ZIP: 94111
TELECOMMUNICATION INFORMATION:
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 2257 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
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Pred. No. 1.3e-236;
                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
-GPGAGAGSGAGAGSGAGAGSGVGVPGVG
                                                                                 -GPGAGAGSGAGAGSGAGAGSGVGV
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RESULT 9
US-08-477-5098-82
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GENERAL INFORMATION:
APPLICANT: Ferrari, Franco A
APPLICANT: Cappello, Joseph
APPLICANT: Crissman, John w
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                                                                                                                                                                                                                                                                    CLASSIFICATION: 435

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/175,155
FILING DATE: 29-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/053,049
FILING DATE: 22-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/114,618
FILING DATE: 29-OCT-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 06/927,258
                                                                                                                           TELEFAX: 415-398-3249 INFORMATION FOR SEQ ID NO:
                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                 REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-55186-7/RFT/MTK
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Crissman, John w
APPLICANT: Dorman, Mary A
TITLE OF INVENTION: No. 5770697el Peptides Comprising Repetitive
TITLE OF INVENTION: Units of Amino Acids and DNA Sequences Encoding
MOLECULE TYPE:
                      TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/477,509B FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: Four Embarc:
CITY: San Francisco
STATE: California
                                                            TYPE: amino acid
                                                                                                                                                                                                                                                        NAME: Trecartin, Richard F.
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                                                                                  LENGTH:
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                                     VPGVGVPGVGVPGVGVPGVGVPGVGVPGAGAGSGAGAGSGAGAGSGAGAGSGAGA
                                                       VPGVGVPGKGVPGVG-PGVG-PGV------
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Pred. No. 1.3e-236;
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RESULT 10
US-08-707-237A-53
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                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 07/609,716
FILING DATE: 06-NOV-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/269,429
FILING DATE: 09-NOV-1988
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/114,618
APPLICATION NUMBER: US 07/114,618
                                                                                    TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 2257 amino acids
                                                                                                                                                                                                                     FILING DATE: 04-NOV-1986
ATTORNEY/AGENT INFORMATION:
NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-55
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 94111-4187

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Capello, Joseph
APPLICANT: Crissman, John W.
APPLICANT: Dorman, Mary A.
TITLE OF INVENTION: METHODS FOR PREPARING SYNTHETIC
TITLE OF INVENTION: REPETITIVE DNA
NUMBER OF SEQUENCES: 108
  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 29-OCT-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 22-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/609,716
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/175,155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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CITY: San Francisco
STATE: California
COUNTRY: United States
                         TOPOLOGY:
                                           TYPE: amino acid
STRANDEDNESS: uni
                                                                                                                                                                              TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
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-GPGAGAGSGAGAGSGAGAGSGVGVPGVGVPGV 744
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RESULT 11
US-08-482-085B-82
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                                                                              US-08-482-085B-82
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            Matches
                         Query Match
Best Local
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FILING DATE: 29-OCT-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,049
FILING DATE: 22-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/175,155
FILING DATE: 29-DEC-1993
                                                                                                                                                                                  TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TOPOLOGY: lin
MOLECULE TYPE:
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APPLICANT:
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APPLICANT: Cappello, Joseph
APPLICANT: Crissman, John W.
TITLE OF INVENTION: No. 6018030el Peptides Comprising Repetitive
TITLE OF INVENTION: Units of Amino Acids and DNA Sequences Encoding
NUMBER OF SEQUENCES: 112
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                                               REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-55186-6/RFT/MTK
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415.781-1989
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APPLICATION NUMBER: US 00
FILING DATE: 04-NOV-1986
PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US,
FILING DATE: 07-JUN-19:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                        FILING DATE: 29-DEC-1993
ATTORNEY/AGENT INFORMATION:
Local 5.755;
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CITY: San Francis
STATE: California
                                                                                                                                               TYPE: amino acid
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                                                                                                                                                                                                                                                                                                        NAME: Trecartin, Richard F
                                                                                                                                                                 LENGTH:
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Chambers, James
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         Score 3296; DB 2;
Pred. No. 1.3e-236;
0; Mismatches 25;
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Suite 3400
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; SEQUENCE DESCRIPTION: SEQ ID NO: 82:
US-09-444-791A-82
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US-09-444-791A-82
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                                                                                                                                           Query Match
Best Local Similarity
Matches 755; Conserv
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INFORMATION FOR SEQ ID NO: 82:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/482,085
FILING DATE: 07-JUN-1995
APPLICATION NUMBER: US 08/175,155
FILING DATE: 29-DEC-1993
APPLICATION NUMBER: US 08/053,049
FILING DATE: 22-DER-1993
APPLICATION NUMBER: US 07/114,618
FILING DATE: 29-OCT-1987
APPLICATION NUMBER: US 06/927,258
FILING DATE: 04-NOV-1986
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-55186-11/RFT/BTC
TELECOMMUNICATION INFORMATION:
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FILING DATE: 22-No. 6355776-1999
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr Hohbach Test Albritton & Herbert
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86
                                    47
                                                                      38
GAGAGSGAGAGSGVGVPGVGVPGVGVPGVG-PGVG-PGVG-PGVG-PGV-------
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CITY: San Francisco
STATE: California
COUNTRY: US
                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                 LENGTH: 2257 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Trecartin, Richard F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cappello, Joseph
Crissman, John W.
OF INVENTION: No. 6355776el Peptides Comprising Repetitive
Units of Amino Acids and DNA Sequences Encoding the
                                                                                                                                                 Conservative
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Pollock, Thomas J.
                   79.0%;
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                                                                                                                                             Score 3296; DB 2;
Pred. No. 1.3e-236;
0; Mismatches 25;
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RESULT 13
US-08-397-633A-54
; Sequence 54, Application US/08397633A
; Patent No. 5773577
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; TOPOLOGY: linear
; MOLECULE TYPE: prof
US-08-397-633A-54
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Best Local Similarity
Matches 718; Conserv
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: PATENTIN Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/397,633A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Rowland, Bertram I
REGISTRATION NUMBER: 20,015
REFERENCE/DOCKET NUMBER: A-58848-1/BIR PROP-011-1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 877 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cappello, Joseph
TITLE OF INVENTION: PRODUCTS COMPRISING SUBSTRATESCAPABLE
TITLE OF INVENTION: OF ENZYMATIC CROSS-LINKING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: PLEHR, HOR
STREET: 4 Embarcadero
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94111-4187
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                E: FLEHR, HOHBACH, TEST, ALBRITTON 4 Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                  protein
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                                                                                                                                                                                                                                                                                                                                                                      Score 3284; DB 1;
Pred. No. 4.5e-236;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Version
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                                                                                                              ; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-212-237-4
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US-08-212-237-4
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                                                       Matches
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                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.

ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V6
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Cappel
                                                                                                                                                                                                               TELEPHONE: 415-781-1989
TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: CA
                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                           NAME: Rowland, Bertram I
REGISTRATION NUMBER: 20,015
REFERENCE/DOCKET NUMBER: A-58847/BIR
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
TELEPAX: 415-398-3249
                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/212,237
FILING DATE: 11-MAR-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Cappello, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                   TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  717
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                GAGSGVGVPGVGVPGVGVPGKGVPGVG-PGVG-PGVG-PGVGPGAGAGSGAGAGSGAGAG
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Synthetic Proteins As
                                                    Score 3253; DB 1; Length 8 Pred. No. 8.5e-234; 0; Mismatches 13; Indels
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                                                                                Length 832;
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US-08-806-029-27
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                                                                             Sequence 27, Application US/08806029
Patent No. 6380154
Patent No. 6380154
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Stedronsky, Erwin R.
APPLICANT: Stedronsky, Erwin R.
TITLE OF INVENTION: Synthetic Proteins for in vivo Drug
TITLE OF INVENTION: Delivery and Tissue Augmentation
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                       STREET: Four Embarcade:
CITY: San Francisco
STATE: California
COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                     674
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589

397 412 453 471 471 508 285 294 341 175

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; LENGTH: 832 amino acids
; TYPE: amino acid
; STRANDEDNESS: unknown
; TOPOLOGY: unknown
; MOLECULE TYPE: protein
US-08-806-029-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/212,237
FILING DATE: 11-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: TRECARTIN, Richard F.
REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-58847-2/RFT/MTK
TELEPHONE: (415) 781-1989
TELEPHONE: (415) 398-3249
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 832 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 78.0%;
Best Local Similarity 88.9%;
Matches 742; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRINT APPLICATION DATA:
APPLICATION NUMBER: US/08/806,029
FILING DATE: 24-FEB-1997
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                                                   VGVPGVGVPGVGVPGVGVPGVG-GAGAGSGAGAGSGAGAGSVPGVGVPGVGVPG
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Pred. No. 8.5e-234;
0; Mismatches 13; Indels 8
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Search completed: May 10, 2006, 10:45:25 Job time : 55 BECB

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Result
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Maximum
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     Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being pland is derived by analysis of the total score distribution.
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   length: 0
length: 2000000000
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
   GenCore version 5.1.8 (c) 1993 - 2006 Biocceleration Ltd.
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US-10-441-965-19
US-10-800-179-19
US-10-845-936A-19
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US-10-816-029-15
US-10-816-775A-25
US-10-816-775A-25
US-10-845-936A-32
US-10-845-936A-32
US-10-845-936A-31
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63.3	63.6	63.9	64.1	65.7	67.0	67.3	67.4	67.8	68.4	68.5	71.8	74.0	75.7	76.1	76.3	76.5	77.9
972	936	1011	953	966	936	696	696	750	1002	696	1056	2018	2055	1465	988	768	1043
N	N	4.	N	4.	4	4	N	N	4	4.	N	4	4	4	N	N	u
US-08-806-029-30	US-08-806-029-26	US-10-096-986-94	US-08-806-029-14	US-10-117-931-34	US-10-117-931-30	US-10-441-965-23	US-08-806-029-36	US-08-806-029-25	US-10-117-931-25	US-10-441-965-21	US-08-806-029-29	US-10-096-986-80	US-10-096-986-81	US-10-096-986-74	US-08-806-029-28	US-08-806-029-35	US-10-845-936A-35
Sequence 30	Sequence 26	Sequence 94	Sequence 14	Sequence 34	Sequence 30	Sequence 23	Sequence 36	Sequence 25	Sequence 25	Sequence 21	Sequence 29	Sequence 80	Sequence 81	Sequence 74	Sequence 28	Sequence 35	Sequence 35
Appl	, Appl	, Appl	Appl	•	•	•	•	•	•	•	•	•	-	•	-	Appl	•

ALIGNMENTS

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; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: polymer US-10-441-965-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-441-965-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: SYNTHESIS OF INORGANIC STRUCTURES USING TEMP TITLE OF INVENTION: CATALYSIS BY SELF ASSEMBLED REPEAT PROTEIN FILE REFERENCE: DOC 0061 PA CURRENT APPLICATION NUMBER: US/10/441,965 CURRENT FILING DATE: 2003-05-20 PRIOR APPLICATION NUMBER: 60/381,913 PRIOR APPLICATION NUMBER: 60/381,913 PRIOR FILING DATE: 2002-05-20 PRIOR PRIOR DATE: 2002-05-20 PRIOR PRIOR DATE: 2002-05-20 PRIOR PRIOR PRIOR DATE: 2002-05-20 PRIOR PRIOR PRIOR DATE: 2002-05-20 PRIOR DATE: PACENTIN Ver. 2.1
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Publication No. US20040014186A1
GENERAL INFORMATION:
APPLICANT: KUMAR, MANOJ
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Best Local Similarity
Matches 780; Conserv
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ORGANISM: Artificial Sequence
FEATURE:
                             301
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                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cuevas, William A.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj
TITLE OF INVENTION: Use of Repeat Sequence Protein Polymers in Personal Care
TITLE OF INVENTION: Compositions
FILE REFERENCE: DOC 0057 PA / GC792-4 / DC 5074
CURRENT FILING DATE: 2004-03-12
CURRENT FILING DATE: 2004-03-12
PRIOR APPLICATION NUMBER: 60/454,077
PRIOR FILING DATE: 2003-03-12
NUMBER OF SEQ ID NOS: 31
SOFTWARE: Patentin version 3.2
SEQ ID NO 19
LENGTH: 780
                                                                                                                                                                                                                  Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19, Application US/10800179 Publication No. US20040180027A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Unknown
FEATURE:
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US-10-845-775A-19

VS-10-845-775A-19

i Sequence 19, Application US/10845775A

publication No. US20040228913A1

i GENERAL INFORMATION:
   APPLICANT: Kumar, Manod
   APPLICANT: Mazeaud, Isabelle
   APPLICANT: Christiano, Steven P.
   TITLE OF INVENTION: Controlled Release of Active Agents Utilizing Repeat Sequence
   TITLE OF INVENTION: Protein Polymers
   FILE REFERENCE: DOC 0077 PA/GC 792-6/DC 5110
   CURRENT FILING DATE: 2004-05-14
   PRIOR APPLICATION NUMBER: US/10/845,775A

CURRENT FILING DATE: 2003-05-14

NUMBER OF SEQ ID NOS: 31

SOPTWARE: PatentIn version 3.2

SEQ ID NO 19

LENGTH: 780
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; OTHER INFORMATION: silk and elastin peptide repeats
US-10-845-775A-19
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Best Local Similarity
Matches 780; Conserv
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                                                                                                    Conservative
                                                                                                100.0%; Score 4173; DB 5;
100.0%; Pred. No. 6.6e-248;
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APPLICANT: COLLIER, Katherine D.
APPLICANT: Cuevas, William A.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: Repeat Sequence Protein Pol
TITLE OF INVENTION: and Uses
FILE REFERENCE: DOC0068pA,DC5058/GC792
FILE REFERENCE: DOC0068pA,DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 37
SOPTWARE: PatentIn version 3.2
SEQ ID NO 19
LENGTH: 780
TYPE: PRT
ORGANISM: unknown
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                                                                                                                                                                                                       Sequence 19, Application US/10845936A Publication No. US20040234609A1 GENERAL INFORMATION:
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US-10-939-036-19

; Sequence 19, Application US/10939036

; Publication No. US20050142094A1

; GENERAL INFORMATION:
APPLICANT: Kumar, Manoj
TITLE OF INVENTION: Use of Repeat Sequence Protein Pol
TITLE OF INVENTION: in Personal Care Compositions
FILE REFERENCE: GC792-4-2
CURRENT APPLICATION NUMBER: US/10/939,036
CURRENT APPLICATION NUMBER: US/10/939,036
CURRENT APPLICATION NUMBER: US 10/800,179
PRIOR APPLICATION NUMBER: US 60/454,077
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RESULT 6
US-08-806-029-19
; Sequence 19, Application US/08806029
; Publication No. US20020045567A1
; GENERAL INFORMATION:
APPLICANT: Cappello, Joseph
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SEQ ID NO 19
LENGTH: 780
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; TYPE: amino acid
; STRANDEDNESS: unkno
; TOPOLOGY: unknown
; MOLECULE TYPE: protei
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Best Local Similarity
Matches 765; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-58847-2/RFT/MTK
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: FOUL CITY: San Francisco
STATE: California
"""ired State
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Synthetic Proteins for in vivo Dru
TITLE OF INVENTION: Delivery and Tissue Augmentation
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 889 amino acids
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APPLICATION NUMBER: US 0
FILING DATE: 11-MAR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, CURRENT APPLICATION DATA:
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Pred. No. 2.6e-208;
0; Mismatches 13;
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Gaps

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SEQUENCE DESCRIPTION: SEQUENCE DESCRIPTION: SEQUENCE DESCRIPTION: S
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Publication No. US20030104589A1
GENERAL INFORMATION:
  Query Match
                                                                                                            TELEFAX: 415-398-3249 INFORMATION FOR SEQ ID NO: 15: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/117,931
FILING DATE: 05-Apr-2002
CLASSIFICATION: «Unknown»
                                                                                                                                             PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/642,246

FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:

NAME: ROWLAND, Bertram I

REGISTRATION NUMBER: 20015

REGISTRATION NUMBER: A61127-1/BIR

TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST,
STREET: Four Embarcadero Center,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: STEDRONSKY, Erwin R.
CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive Using Synthetic
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                                                 TOPOLOGY: linear MOLECULE TYPE: peptid
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                                                                        STRANDEDNESS: single
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                                      SEQ
  Score 3448;
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  Length 884;
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US-10-800-179-25
; Sequence 25, Application US/10800179
; Publication No. US20040180027A1
; GENERAL INFORMATION:
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APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj
TITLE OF INVENTION: Use of Repeat Sequence Protein
TITLE OF INVENTION: Compositions
FILE REFERENCE: DOC 0057 PA / GC792-4 / DC 5074
CURRENT APPLICATION NUMBER: US/10/800,179
CURRENT FILING DATE: 2004-03-12
PRIOR APPLICATION NUMBER: 60/454,077
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NUMBER OF SEQ ID NOS: 31
SOFTWARE: PatentIn version 3.2
SEQ ID NO 25
LENGTH: 884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 82.6%;
Best Local Similarity 90.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Unknown
FEATURE:
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Pred. No. 1.5e-203;
0; Mismatches 26;
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RESULT

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Sequence 25, Application US/10845775A
publication No. US20040228913A1
GENERAL INFORMATION:
APPLICANT: Kumar, Manoj
APPLICANT: Mazeaud, Isabelle
APPLICANT: Mazeaud, Isabelle
APPLICANT: Mazeaud, Isabelle
TITLE OF INVENTION: Controlled Release of Active Age
TITLE OF INVENTION: Protein Polymers
FILE REFERENCE: DOC 0077 PA/GC 792-6/DC 5110
CURRENT APPLICATION NUMBER: US/10/845,775A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: US 60/470,465
PRIOR FILING DATE: 2003-05-14
VINDBER OF SEQ ID NOS: 31
SOFTWARE: Patentin version 3.2
SEQ ID NO 25
LENGTH: 884
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                               PGXGVPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVP
                                                                           VPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPGVGVPGVGV
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Pred. No. 1.5e-203;
0; Mismatches 26;
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FILE REFERENCE: DOCOGERPA/DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
INUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.2
SEQ ID NO 25
LENGTH: 884
TYPE: PRT
ORGANISM: unknown
FEATURE:
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US-10-845-936A-25
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US-10-845-936A-25
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Publication No. US20040234609A1
GENERAL INFORMATION:
APPLICANT: Collier, Katherine D.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: Repeat Sequence Protein
TITLE OF INVENTION: and Uses
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Matches
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Local Similarity 90.6%;
hes 752; Conservative
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Pred. No. 1.5e-203;
O; Mismatches 26;
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TITLE OF INVENTION: in Personal Care Compositions
FILE REFERENCE: GC792-4-2
CURRENT APPLICATION NUMBER: US/10/939,036
CURRENT FILING DATE: 2004-09-10
PRIOR APPLICATION NUMBER: US 10/800,179
PRIOR FILING DATE: 2004-03-12
PRIOR FILING DATE: 2004-03-12
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2003-03-12
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 25
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US-10-939-036-25
; Sequence 25, Application US/10939036
; Publication No. US20050142094A1
; GENERAL IMPORMATION:
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                                                                                                                                                                                                                                                                 Query Match 82.6%;
Best Local Similarity 90.6%;
Matches 752; Conservative
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                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Unknown
FEATURE:
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Pred. No. 1.5e-203;
0; Mismatches 26;
                                                                                                                                                                                                                                                                                            Length 884;
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; OTHER INFORMATION: ; OTHER INFORMATION: US-10-845-936A-37
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US-10-845-936A-37
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Publication No. US20040234609A1
GENERAL INFORMATION:
APPLICANT: Collier, Katherine D.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: Repeat Sequence Protein Polymer .
TITLE OF INVENTION: and Uses
FILE REFERENCE: DOC0068PA/DC5058/GC792
                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 37
SOFTWARE: PatentIn version 3.2
SEQ ID NO 37
TENOTH: 983
                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                            FEATURE:
                                                                                                                                                                                      ORGANISM: unknown
                                                                                                                                                                                                        TYPE: PRT
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                                                                                             90.6%;
                                                                              Score 3448; DB 5;
Pred. No. 1.6e-203;
0; Mismatches 26;
                                                                                  Indels
                                                                                                          Length
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                                    Sequence 32, Application US/10845936A
Publication No. US20040234609A1
GENERAL INFORMATION:
APPLICANT: COllier, Katherine D.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: Repeat Sequence Protein Pol
TITLE OF INVENTION: and Uses
FILE REFERENCE: DOC0068PA/DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT ETLING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.2
SEQ ID NO 32
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LENGTH: 1027
TYPE: PRT
ORGANISM: unknown
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RESULT 14
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US-10-845-936A-33
; Sequence 33, Application US/10845936A
; Publication No. US20040234609A1
; Publication No. US20040234609A1
; GENERAL INFORMATION:
; APPLICANT: Collier, Katherine D.
; APPLICANT: Cuevas, William A.
; APPLICANT: Kumar, Manoj A.
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Pred. No. 1.7e-203;
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FITTLE OF INVENTION: Repeat Sequence Protein Pol
TITLE OF INVENTION: and Uses
FILE REFERENCE: DOCOGO6BA/DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 37
SOPTWARE: PatentIn version 3.2
SEQ ID NO 33
LENGTH: 1105
TYPE: PRT
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: US-10-845-936A-33
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Best Local Simi
Matches 752;
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Pred. No. 1.8e-203;
D; Mismatches 26;
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Best Local S
Matches 752
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Publication No. US20040234609A1
GENERAL INFORMATION:
APPLICANT: Collier, Katherine D.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: Repeat Sequence Protein Polymer Active Agent Conjugates, Methods
TITLE OF INVENTION: and Uses
TILE REFERENCE: DOC0068PA/DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/845,936A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION UNMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Datentt "Maria" 3 2
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TYPE: PRT
ORGANISM: unknown
FEATURE:
OTHER INFORMATION: GFP-SELP47K: silk, elastin and green fluorescent protein peptides
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90.6%; Pred. No. 1.8e-203;
tive 0; Mismatches 26; Indels 52
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         GVPGVGVPGVGVPGKGVPGVG-PGVG-PGVG-PGVG-PGAGAGSGAGAGS
                                                                                   GVGVPGKGVPGVG-PGVG-PGVG-PGAGAGSGAGAGSGAGAGSGAGAGSGVGVPG
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                                   VGVPGVGVPGVGVPGVGVPGVGVPGVGVPGAGAGSGAGAGSGAGAGSGAGAGSGV
                  780
                                     1054
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Search completed: May 10, 2006, Job time: 171 secs 10:48:14

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Result
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                       Published Applications AA New:*

1: /SIDSS/prodata/2/pubpaa/USO8 NEW_PUB.pep1:*

2: /SIDSS/prodata/2/pubpaa/USO6 NEW_PUB.pep:*

3: /SIDSS/prodata/2/pubpaa/USO7 NEW_PUB.pep:*

4: /SIDSS/prodata/2/pubpaa/USO8 NEW_PUB.pep:*

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6: /SIDSS/prodata/2/pubpaa/USO9 NEW_PUB.pep:*

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11: /SIDSS/prodata/2/pubpaa/USO1 NEW_PUB.pep:*

12: /SIDSS/prodata/2/pubpaa/USO1 NEW_PUB.pep:*
                                                                                                                                                                                                                                                                                                                                           Match
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                              Minimum Match 0%
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Listing first 45 summaries
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RESULT 1
US-11-201-606-15
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24.5	24.5		26.1	26.3	26.5	26.5	26.5	26.5	26.8	26.8	31.4	32.8	33.8	•	•	33.9		34.0	34.1		34.2	34.2	34.2
728	624	1660	1079	1901	357	356	354	339	357	574	378	450	879	775	507	506	504	489	859	864	774	757	736
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US-11-053-100-54	US-11-053-100-48	US-11-052-554A-137	US-11-052-554A-145	US-11-052-554A-135	US-11-053-100-33	US-11-053-100-36	US-11-053-100-27	US-11-053-100-23	US-11-053-100-30	US-11-053-100-40	US-11-201-606-26	US-11-053-100-16	US-11-053-100-57	US-11-053-100-51	US-11-053-100-34	US-11-053-100-37	US-11-053-100-28	US-11-053-100-24	US-11-053-100-52	US-11-053-100-58	US-11-053-100-49	US-11-053-100-47	US-11-053-100-45
54,	48,	Sequence 137, App	Sequence 145, App	Sequence 135, App	ω ω,	36,	27,	•	30,	-	-	16,		•	•		•	•	52,	58,	49,	•	•

ALIGNMENTS

## Sequence 15, Application US/11201606 Publication No. US20060029638A1 STATE: CA COUNTRY: US ZIP: 94111 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC Compatible COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Ver CURRENT APPLICATION DATA: APPLICATION UNMBER: US/11/201,606 FILING DATE: 2005-08-10 CLASSIFICATION PRIOR APPLICATION DATA: APPLICATION NUMBER: US/08/642,246 FILING DATE: 1996-05-02 CLASSIFICATION NUMBER: US/08/642,246 FILING DATE: 1996-05-02 TELEFAX: 415-398-3249 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: CLASSIFICATION: ATTORNEY AGENT INFORMATION: NAME: ROWLAND, Bertram I REGISTRATION NUMBER: 20015 REFERENCE/DOCKET NUMBER: A611 TELECOMMUNICATION INFORMATION: TELEPHONE: 415-781-1989 GENERAL INFORMATION: APPLICANT: STEDRONSKY, Erwin R. APPLICANT: CAPPELLO, Joseph TITLE OF INVENTION: Tissue Adhesive TITLE OF INVENTION: Crosslinking CORRESPONDENCE ADDRESS: NUMBER OF SEQUENCES: MOLECULE LENGTH: 884 a TYPE: amino a STRANDEDNESS: TOPOLOGY: lin STREET: Four Embarca CITY: San Francisco ADDRESSEE: amino acid 884 amino acids E: FLEHR, HOHBACH, TEST, Four Embarcadero Center, linear single 15: A61127-1/BIR ALBRITTON Suite 200 Version Using Synthetic #1.30 & HERBERT

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RESULT 2
US-11-201-606-25
US-11-201-606-25
; Sequence 25, Application US/11201606
; Publication No. US20060029638A1
; Publication WFORMATION:
GENERAL INFORMATION:
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Best Local
APPLICANT: STEDRONSKY, Erwin R.
APPLICANT: CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive Using
TITLE OF INVENTION: Crosslinking
NUMBER OF SEQUENCES: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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Pred. No. 6.6e-230;
0; Mismatches 26;
               Synthetic
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; LENGTH: 1002 amino
; TYPE: amino acid
; STRANDEDNESS: sing
; TOPOLOGY: linear
; MOLECULE TYPE: prote
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Best Local Similarity 65.0
Matches 668; Conservative
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APPLICATION NUMBER: US/08/642,246
FILING DATE: 1996-05-02
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A61127-1/BIR
TELEPHONE: 415-781-199
TELEPHONE: 415-781-199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/201,606
FILING DATE: 2005-08-10
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 415-781-190
TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
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ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON
STREET: Four Embarcadero Center, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 1002 amino acids
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CITY: San Francisco
STATE: CA
COUNTRY: US
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                         linear
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RESULT 3
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IS-11-201-606-30
; Sequence 30, Application US/11201606
; Publication No. US20060029638A1
; GENERAL INFORMATION:
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                       ZIP: 94111

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-DOS/MS-DOS
SOFTWARE: PATENTIN PATA:
APPLICATION NUMBER: US/11/201,606
FILING DATE: 2005-08-10
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/642,246
FILING DATE: 1996-05-02
CLASSIFICATION:
NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A61127-1/BIR
                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST,
STREET: Four Embarcadero Center,
                                                                                                                                                                                                                                                                      APPLICANT: STEDRONSKY, Erwin R.
APPLICANT: CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive
TITLE OF INVENTION: Crosslinking
        TELECOMMUNICATION INFORMATION: TELEPHONE: 415-781-1989
                                                                                                                                                                                                    CITY: San Fr.
STATE: CA
COUNTRY: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: sin
TOPOLOGY: linear
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Pred. No. 4.1e-185;
4; Mismatches 93;
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US-11-201-606-34
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Publication No. US20060029638A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 65.7%;
Best Local Similarity 72.5%;
Matches 642; Conservative
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APPLICANT: CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive Using Sy.
TITLE OF INVENTION: Crosslinking
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
ADDRESSE: FOLERR, HOHBACH, TEST, ALBRITTON
STREET: FOUR Embarcadero Center, Suite 200
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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NAME: ROWLAND, Bertram I
REGISTRAFION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A61127-1/BIR
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
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APPLICATION NUMBER: US/
FILING DATE: 1996-05-02
CLASSIFICATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/201,606
FILLING DATE: 2005-08-10
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
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Pred. No. 2.5e-181;
4; Mismatches 104;
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Suite 200
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Sequence 46, Application US/11053100

Publication No. US2005025554A1

GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY
FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILING DATE: 2001-03-20
PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR FILING DATE: 2000-03-20
NUMBER OF 502 ID NOS: 58
SOFTWARE: PatentIn version 3.3

1.ENGTHAND: 1166
                                                                                                                        ; OTHER INFORMATION: Synthetic Construct
; FEATURE:
; NAME/KEY: MISC FEATURE
; LOCATION: (1). (1186)
; OTHER INFORMATION: pET15b-SD3-ELP1-180-throm-Androgen receptor ligand
US-11-053-100-46
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                                                     Query Match
Best Local S
Matches 542
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TYPE: PRT
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                                                                       Similarity
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                                                        Conservative
                                                                       59.2%;
                                                     Score 2091.5; DB 11;
Pred. No. 8.4e-137;
2; Mismatches 223; II
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                                                        Indels 149; Gaps
                                                                                        Length 1186;
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RESULT 6
US-11-053-100-53
; Sequence 53, Application US/11053100
; Publication No. US2005025554A1
; GENERAL INFORMATION:
; APPLICANT: CHILKOTI, Ashutosh
; TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY
; FILE REFERENCE: 4176-101 CIP
; CURRENT APPLICATION NUMBER: US/11/053,100
; CURRENT FILING DATE: 2005-02-08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09
PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: Patentin version 3.3
SEQ ID NO 53
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ORGANISM: Artificial
FEATURE:
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                    GKGVPGVG-------PGVG--PGVG-PGAGA-GSGAGAGSGAGAGSGAGAGSGVGVVP
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Pred. No. 1.7e-136;
3; Mismatches 223; I
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US-11-053-100-26
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SOFTWARE: Patentin version 3.3
SEQ ID NO 26
LENGTH: 939
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                                                                                                                                                                                                                                                                                                                                                                                                               Matches 539;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE
FILE REFERENCE: 4176-101 CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: MISC_FEATURE
LOCATION: (1)..(939)
OTHER INFORMATION: pET32a-SD15-ELP1-180-EK-Insulin
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ORGANISM: Artificial
FEATURE:
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Pred. No. 1.7e-136;
2; Mismatches 225;
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; OTHER INFORMATION: Synthetic Construct; FEATURE; NAME/KEY: MISC_FEATURE
; LOCATION: (1)...(1174)
; OTHER INFORMATION: pET15b-SD5-ELP1-180-throm-Tobacco etch virus
US-11-053-100-43
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PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
SEQ ID NO 43
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 43, Application US/11053100 Publication No. US20050255554A1 GENERAL INFORMATION:
                                                      Query Match
Best Local Similarity
Matches 539; Conserv
                                                                                                                                                                                                                                                                                                                         APPLICANT: CHILKOTI, AShutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE
FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILLING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILLING DATE: 2001-03-20
                                                                                                                                                                                              ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                             TYPE: PRT
                                                                                                                                                                                                                                        LENGTH: 1174
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                                                    Score 2086; DB 11;
Pred. No. 2e-136;
2; Mismatches 225;
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Sequence 50, Application US/11053100
Publication No. US2005025554A1
GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE
FILE REFERENCE: 4176-101 CDP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
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PRIOR APPLICATION NUMBER: US 60
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
SEQ ID NO 50
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ORGANISM: Artificial
FEATURE:
OTHER INFORMATION: Synthetic Co
FEATURE:
FEATURE:
NAME/KEY: MISC FEATURE
NAME/KEY: MISC TEATURE
TOTATION: (1)...(1225)
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Best Local Similarity
Matches 539; Conserv
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Pred. No. 2.1e-136;
2; Mismatches 225;
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LOCATION: (1). (1225)
OTHER INFORMATION: pET15b-SD5-ELP1-180-throm-Estrogen receptor ligand binding domain
 Indels
                             Length
                                 1225;
140;
Gaps
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FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 200-2-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: Patentin version 3.3
SEQ ID NO 56
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US-11-053-100-56
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Best Local Similarity
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TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY PHASE TRANSITION
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OTHER INFORMATION: pET15b-SD5-ELP1-180-throm-1-Deoxy-D-Xylulose 5-Phosphate OTHER INFORMATION: Reductoisomerase Peptide
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ORGANISM: Artificial
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Pred. No. 2.2e-136;
2; Mismatches 225;
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                                                                  Query Match
Best Local S
Matches 492
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publication No. US2005025554A1
GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
TITLE OF INCENTION: FUSION PEPTIDES ISOLATABLE
                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: Patentin version 3...
SEQ ID NO 32
                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILING DATE: 2001-03-20
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                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial
PEATURE:
OTHER INFORMATION: Synthetic Construct
PEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (1)...(657)
OTHER INFORMATION: pET17b-ELP4-120-Throm-T20 peptide
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                                                                                    Local Similarity
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                                                                      Conservative
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                                                                    Score 1913.5; DB 11;
Pred. No. 8.3e-125;
0; Mismatches 65; I
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APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY
FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT APPLICATION NUMBER: US 09/812,382
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: Patentin version 3.3
SEQ ID NO 25
LENGTH: 639
TYPE: PRT
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US-11-053-100-25
                                                                                                                             Sequence 25, Application US/11053100 Publication No. US20050255554A1 GENERAL INFORMATION:
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   RESULT 13
US-11-053-100-29
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OTHER INFORMATION: PET
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OTHER INFORMATION: Synthetic
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Pred. No. 1.3e-124;
0; Mismatches 66;
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CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 29
LENGTH: 654
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: CHILKOTI, ASHUTOSH
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY PHASE TRANSITION
FILE REFERENCE: 4176-101 CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: MISC_FEATURE LOCATION: (1)..(654)
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Pred. No. 1.3e-124;
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, NAME/KEY: MISC FEATURE
; LOCATION: (1)...(656)
; OTHER INFORMATION: pET17b-ELP4-120-TEV(Q/Y)-T20 peptide
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US-11-053-100-38
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Publication No. US2005025554A1
GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY PHASE TRANSITION
FILE REFERENCE: 4176-101 CTP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 09/812,382
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Best Local S
Matches 490
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NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
SEQ ID NO 38
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ORGANISM: Artificial
FEATURE:
OTHER INFORMATION: Synthetic Construct
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Pred. No. 2.5e-124;
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SEQ ID NO 35
LENGTH: 657
TYPE: PRT
ORGANISM: Artificial
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OTHER INFORMATION: Synthetic Cc
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Best Local S
Matches 490
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Publication No. US2005025554A1
GENERAL INFORMATION:
APPLICANT: CHILKOTI, Ashutosh
TITLE OF INVENTION: FUSION PEPTIDES ISOLATABLE BY PHASE TRANSITION
FILE REFERENCE: 4176-101 CIP
CURRENT APPLICATION NUMBER: US/11/053,100
CURRENT FILING DATE: 2005-02-08
CURRENT FILING DATE: 2005-02-08
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PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/190,659
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 58
SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                         FEATURE:

NAME/KEY: MISC_FEATURE
LOCATION: (1)..(657)

OTHER INFORMATION: pET17b-ELP4-120-TEV(Q/S)-T20 peptide
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Local Similarity 59.8%;
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Score 1906.5; DB 11.
Pred. No. 2.5e-124;
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Search completed: May 10, 2006, 10:50:15 Job time : 118 secs

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